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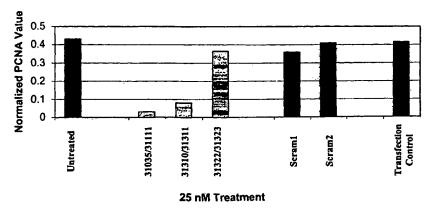
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(54) Title: RNA INTERFERENCE MEDIATED INHIBITION OF GENE EXPRESSION USING SHORT INTERFERING NUCLEIC ACID (SINA)

A549 24h PCNA mRNA Expression



(57) Abstract: The present invention concerns methods and reagents useful in modulating gene expression in a variety of applications, including use in therapeutic, diagnostic, target validation, and genomic discovery applications. Specifically, the invention relates to small nucleic acid molecules, such as short interfering nucleic acid (siNA), short interfering RNA (siRNA), double-stranded RNA (dsRNA), micro-RNA (miRNA), and short hairpin RNA (shRNA) molecules capable of mediating RNA interference (RNAi) against target nucleic acid sequences. The small nucleic acid molecules are useful in the treatment of any disease or condition that responds to modulation of gene expression or activity in a cell, tissue, or organism.

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RNA INTERFERENCE MEDIATED INHIBITION OF GENE EXPRESSION USING SHORT INTERFERING NUCLEIC ACID (siNA)

This invention claims the benefit of Beigelman USSN 60/358,580 filed February 20, 2002, of Beigelman USSN 60/363,124 filed March 11, 2002, of Beigelman USSN 60/386,782 filed June 6, 2002, of Beigelman USSN 60/406,784 filed August 29, 2002, of Beigelman USSN 60/408,378 filed September 5, 2002, of Beigelman USSN 60/409,293 filed September 9, 2002, and of Beigelman USSN 60/440,129 filed January 15, 2003. These applications are hereby incorporated by reference herein in their entireties, including the drawings.

10 Field Of The Invention

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The present invention concerns methods and reagents useful in modulating gene expression in a variety of applications, including use in therapeutic, diagnostic, target validation, and genomic discovery applications. Specifically, the invention relates to small nucleic acid molecules, such as short interfering nucleic acid (siNA), short interfering RNA (siRNA), double-stranded RNA (dsRNA), micro-RNA (miRNA), and short hairpin RNA (shRNA) molecules capable of mediating RNA interference (RNAi).

Background Of The Invention

The following is a discussion of relevant art pertaining to RNAi. The discussion is provided only for understanding of the invention that follows. The summary is not an admission that any of the work described below is prior art to the claimed invention. Applicant demonstrates herein that chemically modified short interfering nucleic acids possess the same capacity to mediate RNAi as do siRNA molecules and are expected to possess improved stability and activity in vivo; therefore, this discussion is not meant to be limiting only to siRNA and can be applied to siNA as a whole.

RNA interference refers to the process of sequence-specific post-transcriptional gene silencing in animals mediated by short interfering RNAs (siRNAs) (Fire *et al.*, 1998, *Nature*, 391, 806). The corresponding process in plants is commonly referred to as post-transcriptional gene silencing or RNA silencing and is also referred to as quelling in fungi. The process of post-transcriptional gene silencing is thought to be an

evolutionarily-conserved cellular defense mechanism used to prevent the expression of foreign genes and is commonly shared by diverse flora and phyla (Fire et al., 1999, Trends Genet., 15, 358). Such protection from foreign gene expression may have evolved in response to the production of double-stranded RNAs (dsRNAs) derived from viral infection or from the random integration of transposon elements into a host genome via a cellular response that specifically destroys homologous single-stranded RNA or viral genomic RNA. The presence of dsRNA in cells triggers the RNAi response though a mechanism that has yet to be fully characterized. This mechanism appears to be different from the interferon response that results from dsRNA-mediated activation of protein kinase PKR and 2',5'-oligoadenylate synthetase resulting in non-specific cleavage of mRNA by ribonuclease L.

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The presence of long dsRNAs in cells stimulates the activity of a ribonuclease III enzyme referred to as dicer. Dicer is involved in the processing of the dsRNA into short pieces of dsRNA known as short interfering RNAs (siRNAs) (Berstein et al., 2001, Nature, 409, 363). Short interfering RNAs derived from dicer activity are typically about 21 to about 23 nucleotides in length and comprise about 19 base pair duplexes (Elbashir et al., 2001, Genes Dev., 15, 188). Dicer has also been implicated in the excision of 21-and 22-nucleotide small temporal RNAs (stRNAs) from precursor RNA of conserved structure that are implicated in translational control (Hutvagner et al., 2001, Science, 293, 834). The RNAi response also features an endonuclease complex, commonly referred to as an RNA-induced silencing complex (RISC), which mediates cleavage of single-stranded RNA having sequence complementary to the antisense strand of the region complementary to the antisense strand of the region complementary to the antisense strand of the siRNA duplex. Cleavage of the target RNA takes place in the middle of the region complementary to the antisense strand of the siRNA duplex (Elbashir et al., 2001, Genes Dev., 15, 188).

RNAi has been studied in a variety of systems. Fire et al., 1998, Nature, 391, 806, were the first to observe RNAi in C. elegans. Wianny and Goetz, 1999, Nature Cell Biol., 2, 70, describe RNAi mediated by dsRNA in mouse embryos. Hammond et al., 2000, Nature, 404, 293, describe RNAi in Drosophila cells transfected with dsRNA. Elbashir et al., 2001, Nature, 411, 494, describe RNAi induced by introduction of duplexes of synthetic 21-nucleotide RNAs in cultured mammalian cells including human embryonic kidney and HeLa cells. Recent work in Drosophila embryonic lysates

(Elbashir et al., 2001, EMBO J., 20, 6877) has revealed certain requirements for siRNA length, structure, chemical composition, and sequence that are essential to mediate efficient RNAi activity. These studies have shown that 21-nucleotide siRNA duplexes are most active when containing 3'-terminal dinucleotide overhangs. Furthermore, complete substitution of one or both siRNA strands with 2'-deoxy (2'-H) or 2'-O-methyl nucleotides abolishes RNAi activity, whereas substitution of the 3'-terminal siRNA overhang nucleotides with 2'-deoxy nucleotides (2'-H) was shown to be tolerated. Single mismatch sequences in the center of the siRNA duplex were also shown to abolish RNAi activity. In addition, these studies also indicate that the position of the cleavage site in the target RNA is defined by the 5'-end of the siRNA guide sequence rather than the 3'-end of the guide sequence (Elbashir et al., 2001, EMBO J., 20, 6877). Other studies have indicated that a 5'-phosphate on the target-complementary strand of a siRNA duplex is required for siRNA activity and that ATP is utilized to maintain the 5'-phosphate moiety on the siRNA (Nykanen et al., 2001, Cell, 107, 309).

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Studies have shown that replacing the 3'-terminal nucleotide overhanging segments a 21-mer siRNA duplex having two -nucleotide 3'-overhangs with deoxyribonucleotides does not have an adverse effect on RNAi activity. Replacing up to four nucleotides on each end of the siRNA with deoxyribonucleotides has been reported to be well tolerated, whereas complete substitution with deoxyribonucleotides results in no RNAi activity (Elbashir et al., 2001, EMBO J., 20, 6877). In addition, Elbashir et al., supra, also report that substitution of siRNA with 2'-O-methyl nucleotides completely abolishes RNAi activity. Li et al., International PCT Publication No. WO 00/44914, and Beach et al., International PCT Publication No. WO 01/68836 preliminarily suggest that siRNA may include modifications to either the phosphate-sugar backbone or the nucleoside to include at least one of a nitrogen or sulfur heteroatom, however, neither application postulates to what extent such modifications would be tolerated in siRNA molecules, nor provides any further guidance or examples of such modified siRNA. Kreutzer et al., Canadian Patent Application No. 2,359,180, also describe certain chemical modifications for use in dsRNA constructs in order to counteract activation of double-stranded RNA-dependent protein kinase PKR, specifically 2'-amino or 2'-Omethyl nucleotides, and nucleotides containing a 2'-O or 4'-C methylene bridge.

However, Kreutzer et al. similarly fails to provide examples or guidance as to what extent these modifications would be tolerated in siRNA molecules.

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Parrish et al., 2000, Molecular Cell, 6, 1977-1087, tested certain chemical modifications targeting the unc-22 gene in C. elegans using long (>25 nt) siRNA transcripts. The authors describe the introduction of thiophosphate residues into these siRNA transcripts by incorporating thiophosphate nucleotide analogs with T7 and T3 RNA polymerase and observed that RNAs with two phosphorothioate modified bases also had substantial decreases in effectiveness as RNAi. Further, Parrish et al. reported that phosphorothioate modification of more than two residues greatly destabilized the RNAs in vitro such that interference activities could not be assayed. Id. at 1081. The authors also tested certain modifications at the 2'-position of the nucleotide sugar in the long siRNA transcripts and found that substituting deoxynucleotides for ribonucleotides produced a substantial decrease in interference activity, especially in the case of Uridine to Thymidine and/or Cytidine to deoxy-Cytidine substitutions. Id. In addition, the authors tested certain base modifications, including substituting, in sense and antisense strands of the siRNA, 4-thiouracil, 5-bromouracil, 5-iodouracil, and 3-(aminoallyl)uracil for uracil, and inosine for guanosine. Whereas 4-thiouracil and 5-bromouracil substitution appeared to be tolerated. Parrish reported that inosine produced a substantial decrease in interference activity when incorporated in either strand. Parrish also reported that incorporation of 5-iodouracil and 3-(aminoallyl)uracil in the antisense strand resulted in a substantial decrease in RNAi activity as well.

The use of longer dsRNA has been described. For example, Beach et al., International PCT Publication No. WO 01/68836, describes specific methods for attenuating gene expression using endogenously-derived dsRNA. Tuschl et al., International PCT Publication No. WO 01/75164, describe a Drosophila in vitro RNAi system and the use of specific siRNA molecules for certain functional genomic and certain therapeutic applications; although Tuschl, 2001, Chem. Biochem., 2, 239-245, doubts that RNAi can be used to cure genetic diseases or viral infection due to the danger of activating interferon response. Li et al., International PCT Publication No. WO 00/44914, describe the use of specific dsRNAs for attenuating the expression of certain target genes. Zernicka-Goetz et al., International PCT Publication No. WO 01/36646, describe certain methods for inhibiting the expression of particular genes in mammalian

cells using certain dsRNA molecules. Fire et al., International PCT Publication No. WO 99/32619, describe particular methods for introducing certain dsRNA molecules into cells for use in inhibiting gene expression. Plaetinck et al., International PCT Publication No. WO 00/01846, describe certain methods for identifying specific genes responsible for conferring a particular phenotype in a cell using specific dsRNA molecules. Mello et al., International PCT Publication No. WO 01/29058, describe the identification of specific genes involved in dsRNA-mediated RNAi. Deschamps Depaillette et al., International PCT Publication No. WO 99/07409, describe specific compositions consisting of particular dsRNA molecules combined with certain anti-viral agents. Waterhouse et al., International PCT Publication No. 99/53050, describe certain methods for decreasing the phenotypic expression of a nucleic acid in plant cells using certain dsRNAs. Driscoll et al., International PCT Publication No. WO 01/49844, describe specific DNA constructs for use in facilitating gene silencing in targeted organisms.

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Others have reported on various RNAi and gene-silencing systems. For example, Parrish et al., 2000, Molecular Cell, 6, 1977-1087, describe specific chemically-modified siRNA constructs targeting the unc-22 gene of C. elegans. Grossniklaus, International PCT Publication No. WO 01/38551, describes certain methods for regulating polycomb gene expression in plants using certain dsRNAs. Churikov et al., International PCT Publication No. WO 01/42443, describe certain methods for modifying genetic characteristics of an organism using certain dsRNAs. Cogoni et al., International PCT Publication No. WO 01/53475, describe certain methods for isolating a Neurospora silencing gene and uses thereof. Reed et al., International PCT Publication No. WO 01/68836, describe certain methods for gene silencing in plants. International PCT Publication No. WO 01/70944, describe certain methods of drug screening using transgenic nematodes as Parkinson's Disease models using certain dsRNAs. Deak et al., International PCT Publication No. WO 01/72774, describe certain Drosophila-derived gene products that may be related to RNAi in Drosophila. Arndt et al., International PCT Publication No. WO 01/92513 describe certain methods for mediating gene suppression by using factors that enhance RNAi. Tuschl et al., International PCT Publication No. WO 02/44321, describe certain synthetic siRNA constructs. Pachuk et al., International PCT Publication No. WO 00/63364, and Satishchandran et al., International PCT Publication No. WO 01/04313, describe certain

methods and compositions for inhibiting the function of certain polynucleotide sequences using certain dsRNAs. Echeverri et al., International PCT Publication No. WO 02/38805, describe certain C. elegans genes identified via RNAi. Kreutzer et al., International PCT Publications Nos. WO 02/055692, WO 02/055693, and EP 1144623 B1 describes certain methods for inhibiting gene expression using RNAi. Graham et al., International PCT Publications Nos. WO 99/49029 and WO 01/70949, and AU 4037501 describe certain vector expressed siRNA molecules. Fire et al., US 6,506,559, describe certain methods for inhibiting gene expression in vitro using certain long dsRNA (greater than 25 nucleotide) constructs that mediate RNAi.

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SUMMARY OF THE INVENTION

This invention relates to compounds, compositions, and methods useful for modulating RNA function and/or gene expression in a cell. Specifically, the instant invention features synthetic small nucleic acid molecules, such as short interfering nucleic acid (siNA), short interfering RNA (siRNA), double-stranded RNA (dsRNA), micro-RNA (miRNA), and short hairpin RNA (shRNA) molecules capable of modulating gene expression in cells by RNA inference (RNAi). The siRNA of the instant invention can be chemically synthesized, expressed from a vector or enzymatically synthesized. The use of chemically modified siNA can improve various properties of native siRNA molecules through increased resistance to nuclease degradation *in vivo* and/or improved cellular uptake. The chemically modified siNA molecules of the instant invention provide useful reagents and methods for a variety of therapeutic, diagnostic, agricultural, target validation, genomic discovery, genetic engineering and pharmacogenomic applications.

In a non-limiting example, the introduction of chemically modified nucleotides into nucleic acid molecules provides a powerful tool in overcoming potential limitations of in vivo stability and bioavailability inherent to native RNA molecules that are delivered exogenously. For example, the use of chemically modified nucleic acid molecules can enable a lower dose of a particular nucleic acid molecule for a given therapeutic effect since chemically modified nucleic acid molecules tend to have a longer half-life in serum. Furthermore, certain chemical modifications can improve the bioavailability of nucleic acid molecules by targeting particular cells or tissues and/or improving cellular uptake of the nucleic acid molecule. Therefore, even if the activity of a chemically modified

nucleic acid molecule is reduced as compared to a native nucleic acid molecule, for example when compared to an all RNA nucleic acid molecule, the overall activity of the modified nucleic acid molecule can be greater than the native molecule due to improved stability and/or delivery of the molecule. Unlike native unmodified siRNA, chemically modified siNA can also minimize the possibility of activating interferon activity in humans.

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The siRNA molecules of the invention can be designed to inhibit gene expression through RNAi targeting of a variety of RNA molecules. In one embodiment, the siRNA molecules of the invention are used to target various RNAs corresponding to a target gene. Non-limiting examples of such RNAs include messenger RNA (mRNA), alternate RNA splice variants of target gene(s), post-transcriptionally modified RNA of target gene(s), pre-mRNA of target gene(s). If alternate splicing produces a family of transcipts that are distinguished by usage of appropriate exons, the instant invention can be used to inhibit gene expression through the appropriate exons to specifically inhibit or to distinguish among the functions of gene family members. For example, a protein that contains an alternatively spliced transmembrane domain can be expressed in both membrane bound and secreted forms. Use of the invention to target the exon containing the transmembrane domain can be used to determine the functional consequences of pharmaceutical targeting of membrane bound as opposed to the secreted form of the protein. Non-limiting examples of applications of the invention relating to targeting these RNA molecules include therapeutic pharmaceutical applications, pharmaceutical discovery applications, molecular diagnostic and gene function applications, and gene mapping, for example using single nucleotide polymorphism mapping with siRNA molecules of the invention. Such applications can be implemented using known gene sequences or from partial sequences available from an expressed sequence tag (EST).

In another embodiment, the siRNA molecules of the invention are used to target conserved sequences corresponding to a gene family or gene families. As such, siRNA can be used to characterize pathways of gene function in a variety of applications. For example, the present invention can be used to inhibit the activity of target gene(s) in a pathway to determine the function of uncharacterized gene(s) in gene function analysis, mRNA function analysis, or translational analysis. The invention can be used to determine potential target gene pathways involved in various diseases and conditions

toward pharmaceutical development. The invention can be used to understand pathways of gene expression involved in development, such as prenatal development, postnatal development and/or aging.

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In one embodiment, the invention features a short interfering nucleic acid (siNA) molecule that down-regulates expression of a gene family by RNA interference. The gene family can comprise more than one splice variant of a target gene, more than one post-transcriptionally modified RNA of a target gene, or more than one RNA trascript having shared homology. In one embodiment, the gene family comprises epidermal growth factor (e.g., EGFR, such as HER1, HER2, HER3, and/or HER4) genes, vascular endothelial growth factor and vascular endothelial growth factor receptor (e.g., VEGF, VEGFR1, VEGFR2, or VEGFR3) genes, or viral genes corresponding to different viral strains (e.g., HIV-1 and HIV-2). Such gene families can be established by analysing nucleic acid sequences (e.g., sequences shown by Genbank Accession Nos. in Table V) for homology.

In one embodiment, the invention features a double-stranded short interfering nucleic acid (siNA) molecule that down-regulates expression of an endogenous mammalian target gene (e.g., a human gene), wherein the siNA molecule comprises one or more chemical modifications and each strand of the double-stranded siNA is about 21 nucleotides long.

In one embodiment, a siNA molecule of the invention comprises no ribonucleotides. In another embodiment, a siNA molecule of the invention comprises ribonucleotides.

In one embodiment, the invention features a double-stranded short interfering nucleic acid (siNA) molecule that down-regulates expression of an endogenous mammalian target gene (e.g., a human gene), wherein one of the strands of the double-stranded siNA molecule comprises a nucleotide sequence that is complementary to a nucleotide sequence of the endogenous mammalian target gene or a portion thereof, and wherein the second strand of the double-stranded siNA molecule comprises a nucleotide sequence substantially similar to the nucleotide sequence of the endogenous mammalian target gene or a portion thereof.

In one embodiment, the invention features a double-stranded short interfering nucleic acid (siNA) molecule that down-regulates expression of an endogenous mammalian target gene (e.g., a human gene), wherein each strand of the siNA molecule comprises about 19 to about 23 nucleotides, and wherein each strand comprises about 19 nucleotides that are complementary to the nucleotides of the other strand.

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In one embodiment, the invention features a double-stranded short interfering nucleic acid (siNA) molecule that down-regulates expression of an endogenous mammalian target gene (e.g., a human gene), wherein the siNA molecule comprises an antisense region comprising a nucleotide sequence that is complementary to a nucleotide sequence of the endogenous mammalian target gene or a portion thereof, and wherein the siNA further comprises a sense region, wherein the sense region comprises a nucleotide sequence substantially similar to the nucleotide sequence of the endogenous mammalian target gene or a portion thereof.

In one embodiment, the invention features a double-stranded short interfering nucleic acid (siNA) molecule that down-regulates expression of an endogenous mammalian target gene (e.g., a human gene), wherein the antisense region and the sense region each comprise about 19 to about 23 nucleotides, and wherein the antisense region comprises about 19 nucleotides that are complementary to nucleotides of the sense region.

In one embodiment, the invention features a double-stranded short interfering nucleic acid (siNA) molecule that down-regulates expression of an endogenous mammalian target gene (e.g., a human gene), wherein the siNA molecule comprises a sense region and an antisense region and wherein the antisense region comprises a nucleotide sequence that is complementary to a nucleotide sequence of RNA encoded by the endogenous mammalian target gene or a portion thereof and the sense region comprises a nucleotide sequence that is complementary to the antisense region.

In one embodiment, the invention features a double-stranded short interfering nucleic acid (siNA) molecule that down-regulates expression of an endogenous mammalian target gene (e.g., a human gene), wherein the siNA molecule is assembled from two separate oligonucleotide fragments wherein one fragment comprises the sense region and the second fragment comprises the antisense region of the siNA molecule.

The sense region can be connected to the antisense region via a linker molecule, such as a polynucleotide linker or a non-nucleotide linker.

In one embodiment, the invention features a double-stranded short interfering nucleic acid (siNA) molecule that down-regulates expression of an endogenous mammalian target gene (e.g., a human gene), wherein the siNA molecule comprises a sense region and an antisense region and wherein the antisense region comprises a nucleotide sequence that is complementary to a nucleotide sequence of RNA encoded by the endogenous mammalian target gene or a portion thereof and the sense region comprises a nucleotide sequence that is complementary to the antisense region, and wherein pyrimidine nucleotides in the sense region are 2'-O-methyl pyrimidine nucleotides, 2'-deoxy nucleotides, and/or 2'-deoxy-2'-fluoro pyrimidine nucleotides.

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In one embodiment, the invention features a double-stranded short interfering nucleic acid (siNA) molecule that down-regulates expression of an endogenous mammalian target gene (e.g., a human gene), wherein the siNA molecule is assembled from two separate oligonucleotide fragments wherein one fragment comprises the sense region and the second fragment comprises the antisense region of the siNA molecule, and wherein the fragment comprising the sense region includes a terminal cap moiety at the 5'-end, the 3'-end, or both of the 5' and 3' ends of the fragment comprising the sense region. In another embodiment, the terminal cap moiety is an inverted deoxy abasic moiety or glyceryl moiety. In another embodiment, each of the two fragments of the siNA molecule comprise 21 nucleotides.

In one embodiment, the invention features a double-stranded short interfering nucleic acid (siNA) molecule that down-regulates expression of an endogenous mammalian target gene (e.g., a human gene), wherein the siNA molecule comprises a sense region and an antisense region and wherein the antisense region comprises a nucleotide sequence that is complementary to a nucleotide sequence of RNA encoded by the endogenous mammalian target gene or a portion thereof and the sense region comprises a nucleotide sequence that is complementary to the antisense region, and wherein the purine nucleotides present in the antisense region comprise 2'-deoxy- purine nucleotides. In another embodiment, the antisense region comprises a phosphorothioate

internucleotide linkage at the 3' end of the antisense region. In another embodiment, the antisense region comprises a glyceryl modification at the 3' end of the antisense region.

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In one embodiment, the invention features a double-stranded short interfering nucleic acid (siNA) molecule that down-regulates expression of an endogenous mammalian target gene (e.g., a human gene), wherein the siNA molecule is assembled from two separate oligonucleotide fragments wherein one fragment comprises the sense region and the second fragment comprises the antisense region of the siNA molecule, and wherein about 19 nucleotides of each fragment of the siNA molecule are base-paired to the complementary nucleotides of the other fragment of the siNA molecule and wherein at least two 3' terminal nucleotides of each fragment of the siNA molecule are not basepaired to the nucleotides of the other fragment of the siNA molecule. In another embodiment, each of the two 3' terminal nucleotides of each fragment of the siNA molecule are 2'-deoxy-pyrimidines, such as 2'-deoxy-thymidine. In another embodiment, all 21 nucleotides of each fragment of the siNA molecule are base-paired to the complementary nucleotides of the other fragment of the siNA molecule. In another embodiment, about 19 nucleotides of the antisense region are base-paired to the nucleotide sequence or a portion thereof of the RNA encoded by the endogenous mammalian target gene. In another embodiment, 21 nucleotides of the antisense region are base-paired to the nucleotide sequence or a portion thereof of the RNA encoded by the endogenous mammalian target gene. In another embodiment, the 5'-end of the fragment comprising said antisense region optionally includes a phosphate group.

In one embodiment, the invention features a double-stranded short interfering nucleic acid (siNA) molecule that inhibits the expression of an endogenous mammalian target RNA sequence (e.g., wherein said target RNA sequence is encoded by a human gene), wherein the siNA molecule comprises no ribonucleotides and wherein each strand of the double-stranded siNA molecule comprises about 21 nucleotides.

In one embodiment, the invention features a double-stranded short interfering nucleic acid (siNA) molecule that inhibits the expression of an endogenous mammalian target gene (e.g., a human gene such as vascular endothelial growth factor, vascular endothelial growth factor receptor (such as VEGFR1, VEGFR2, or VEGFR3), BCL2, HER2/neu, c-Myc, PCNA, REL-A, PTP1B, BACE, CHK1, PKC-alpha, or EGFR),

wherein the siNA molecule does not require the presence of a ribonucleotide within the siNA molecule for said inhibition of expression of an endogenous mammalian target gene and wherein each strand of the double-stranded siNA molecule is about 21 nucleotides long.

In one embodiment, the invention features a medicament comprising a siNA molecule of the invention.

In one embodiment, the invention features an active ingredient comprising a siNA molecule of the invention.

In one embodiment, the invention features the use of a double-stranded short interfering nucleic acid (siNA) molecule to down-regulate expression of an endogenous mammalian target gene, wherein the siNA molecule comprises one or more chemical modifications and each strand of the double-stranded siNA is about 21 nucleotides long.

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In one embodiment, siRNA molecule(s) and/or methods of the invention are used to inhibit the expression of gene(s) that encode RNA referred to by Genbank Accession number in **Table V**. In another embodiment, siRNA molecule(s) and/or methods of the invention are used to target RNA sequence(s) referred to by Genbank Accession number in **Table V**, or nucleic acid sequences encoding such sequences referred to by Genbank Accession number in **Table V**. Such sequences are readily obtained using the Genbank Accession numbers in **Table V**.

In one embodiment, the invention features a siNA molecule having RNAi activity against an RNA encoding a protein, wherein the siNA molecule comprises a sequence complementary to RNA having protein encoding sequence, such as those sequences having GenBank Accession Nos. shown in Table V.

In another embodiment, the invention features a siNA molecule having RNAi activity against a gene, wherein the siNA molecule comprises nucleotide sequence complementary to a nucleotide sequence of the gene, such as genes encoding sequences having GenBank Accession Nos. shown in Table V. In another embodiment, a siNA molecule of the invention includes nucleotide sequence that can interact with nucleotide sequence of a gene and thereby mediate silencing of gene expression, for example,

wherein the siNA mediates regulation of gene expression by cellular processes that modulate the chromatin structure of the gene and prevent transcription of the gene.

In yet another embodiment, the invention features a siNA molecule comprising a sequence, for example, the antisense sequence of the siNA construct, complementary to a sequence represented by GenBank Accession Nos. shown in **Table V** or a portion of said sequence.

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In one embodiment, the nucleic acid molecules of the invention that act as mediators of the RNA interference gene silencing response are chemically modified double stranded nucleic acid molecules. As in their native double stranded RNA counterparts, these siNA molecules typically consist of duplexes containing about 19 base pairs between oligonucleotides comprising about 19 to about 25 nucleotides. The most active siRNA molecules are thought to have such duplexes with overhanging ends of 1-3 nucleotides, for example 21 nucleotide duplexes with 19 base pairs and 2 nucleotide 3'overhangs. These overhanging segments are readily hydrolyzed by endonucleases in vivo. Studies have shown that replacing the 3'-overhanging segments of a 21-mer siRNA duplex having 2 nucleotide 3' overhangs with deoxyribonucleotides does not have an adverse effect on RNAi activity. Replacing up to 4 nucleotides on each end of the siRNA with deoxyribonucleotides has been reported to be well tolerated whereas complete substitution with deoxyribonucleotides results in no RNAi activity (Elbashir et al., 2001, EMBO J., 20, 6877). In addition, Elbashir et al, supra, also report that substitution of siRNA with 2'-O-methyl nucleotides completely abolishes RNAi activity. Li et al., International PCT Publication No. WO 00/44914, and Beach et al., International PCT Publication No. WO 01/68836 both suggest that siRNA may include modifications to either the phosphate-sugar back bone or the nucleoside to include at least one of a nitrogen or sulfur heteroatom, however neither application teaches to what extent these modifications are tolerated in siRNA molecules nor provide any examples of such modified siRNA. Kreutzer and Limmer, Canadian Patent Application No. 2,359,180, also describe certain chemical modifications for use in dsRNA constructs in order to counteract activation of double stranded-RNA-dependent protein kinase PKR, specifically 2'-amino or 2'-O-methyl nucleotides, and nucleotides containing a 2'-O or 4'-C methylene bridge. However, Kreutzer and Limmer similarly fail to show to what

extent these modifications are tolerated in siRNA molecules nor provide any examples of such modified siRNA.

In one embodiment, the invention features chemically modified siNA constructs having specificity for target nucleic acid molecules in a cell (i.e. target nucleic acid molecules comprising or encoded by sequences referred to herein by Genbank Accession numbers in Table V). Non-limiting examples of such chemical modifications include without limitation phosphorothioate internucleotide linkages, 2'-O-methyl ribonucleotides, 2'-deoxy-2'-fluoro ribonucleotides, 2'-deoxy ribonucleotides, "universal base" nucleotides, 5-C-methyl nucleotides, and inverted deoxyabasic residue incorporation. These chemical modifications, when used in various siNA constructs, are shown to preserve RNAi activity in cells while at the same time, dramatically increasing the serum stability of these compounds. Furthermore, contrary to the data published by Parrish et al., supra, applicant demonstrates that multiple (greater than one) phosphorothioate substitutions are well-tolerated and confer substantial increases in serum stability for modified siNA constructs.

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In one embodiment, a siNA molecule of the invention comprises modified nucleotides while maintaining the ability to mediate RNAi. The modified nucleotides can be used to improve in vitro or in vivo characteristics such as stability, activity, and/or bioavailability. For example, a siNA molecule of the invention can comprise modified nucleotides as a percentage of the total number of nucleotides present in the siNA molecule. As such, a siNA molecule of the invention can generally comprise modified nucleotides of about 5 to about 100% of the nucleotide positions (e.g., 5%, 10%, 15%, 20%, 25%, 30%, 35%, 40%, 45%, 50%, 55%, 60%, 65%, 70%, 75%, 80%, 85%, 90%, 95% or 100% of the nucleotide positions). The actual percentage of modified nucleotides present in a given siNA molecule depends on the total number of nucleotides present in the siNA. If the siNA molecule is single stranded, the percent modification can be based upon the total number of nucleotides present in the single stranded siNA molecules. Likewise, if the siNA molecule is double stranded, the percent modification can be based upon the total number of nucleotides present in the sense strand, antisense strand, or both the sense and antisense strands. In addition, the actual percentage of modified nucleotides present in a given siNA molecule can also depend on the total number of purine and pyrimidine nucleotides present in the siNA, for example wherein all

pyrimidine nucleotides and/or all purine nucleotides present in the siNA molecule are modified.

The antisense region of a siNA molecule of the invention can comprise a phosphorothioate internucleotide linkage at the 3'-end of said antisense region. The antisense region can comprise about one to about five phosphorothioate internucleotide linkages at the 5'-end of said antisense region. The 3'-terminal nucleotide overhangs of a siNA molecule of the invention can comprise ribonucleotides or deoxyribonucleotides that are chemically-modified at a nucleic acid sugar, base, or backbone. The 3'-terminal nucleotide overhangs can comprise one or more universal base ribonucleotides. The 3'-terminal nucleotide overhangs can comprise one or more acyclic nucleotides.

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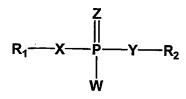
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One embodiment of the invention provides an expression vector comprising a nucleic acid sequence encoding at least one siNA molecule of the invention in a manner that allows expression of the nucleic acid molecule. Another embodiment of the invention provides a mammalian cell comprising such an expression vector. The mammalian cell can be a human cell. The siNA molecule of the expression vector can comprise a sense region and an antisense region. The antisense region can comprise sequence complementary to a RNA or DNA sequence encoding a protein and the sense region can comprise sequence complementary to the antisense region. The siNA molecule can comprise two distinct strands having complementary sense and antisense regions. The siNA molecule can comprise a single strand having complementary sense and antisense regions.

In one embodiment, the invention features a chemically-modified short interfering nucleic acid (siNA) molecule capable of mediating RNA interference (RNAi) inside a cell or reconstituted *in vitro* system, wherein the chemical modification comprises one or more (e.g., about 1, 2, 3, 4, 5, 6, 7, 8, 9, 10, or more) nucleotides comprising a backbone modified internucleotide linkage having Formula I:



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wherein each R1 and R2 is independently any nucleotide, non-nucleotide, or polynucleotide which can be naturally-occurring or chemically-modified, each X and Y is independently O, S, N, alkyl, or substituted alkyl, each Z and W is independently O, S, N, alkyl, substituted alkyl, O-alkyl, S-alkyl, alkaryl, or aralkyl, and wherein W, X, Y, and Z are optionally not all O.

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The chemically-modified internucleotide linkages having Formula I, for example, wherein any Z, W, X, and/or Y independently comprises a sulphur atom, can be present in one or both oligonucleotide strands of the siNA duplex, for example, in the sense strand, the antisense strand, or both strands. The siNA molecules of the invention can comprise one or more (e.g., about 1, 2, 3, 4, 5, 6, 7, 8, 9, 10, or more) chemicallymodified internucleotide linkages having Formula I at the 3'-end, the 5'-end, or both of the 3' and 5'-ends of the sense strand, the antisense strand, or both strands. For example, an exemplary siNA molecule of the invention can comprise about 1 to about 5 or more (e.g., about 1, 2, 3, 4, 5, or more) chemically-modified internucleotide linkages having Formula I at the 5'-end of the sense strand, the antisense strand, or both strands. In another non-limiting example, an exemplary siNA molecule of the invention can comprise one or more (e.g., about 1, 2, 3, 4, 5, 6, 7, 8, 9, 10, or more) pyrimidine nucleotides with chemically-modified internucleotide linkages having Formula I in the sense strand, the antisense strand, or both strands. In yet another non-limiting example, an exemplary siNA molecule of the invention can comprise one or more (e.g., about 1, 2, 3, 4, 5, 6, 7, 8, 9, 10, or more) purine nucleotides with chemically-modified internucleotide linkages having Formula I in the sense strand, the antisense strand, or both strands. In another embodiment, a siNA molecule of the invention having internucleotide linkage(s) of Formula I also comprises a chemically-modified nucleotide or nonnucleotide having any of Formulae I-VII.

In one embodiment, the invention features a chemically-modified short interfering nucleic acid (siNA) molecule capable of mediating RNA interference (RNAi) inside a cell or reconstituted *in vitro* system, wherein the chemical modification comprises one or more (e.g., about 1, 2, 3, 4, 5, 6, 7, 8, 9, 10, or more) nucleotides or non-nucleotides having Formula II:

wherein each R3, R4, R5, R6, R7, R8, R10, R11 and R12 is independently H, OH, alkyl, substituted alkyl, alkaryl or aralkyl, F, Cl, Br, CN, CF3, OCF3, OCN, O-alkyl, S-alkyl, N-alkyl, O-alkyl, S-alkyl, N-alkyl, O-alkyl, S-alkyl, N-alkyl-OH, O-alkyl-OH, O-alkyl-OH, S-alkyl-SH, S-alkyl-SH, alkyl-S-alkyl, alkyl-O-alkyl, ONO2, NO2, N3, NH2, aminoalkyl, aminoacid, aminoacyl, ONH2, O-aminoalkyl, O-aminoacid, O-aminoacyl, heterocycloalkyl, heterocycloalkaryl, aminoalkylamino, polyalklylamino, substituted silyl, or group having Formula I; R9 is O, S, CH2, S=O, CHF, or CF2, and B is a nucleosidic base such as adenine, guanine, uracil, cytosine, thymine, 2-aminoadenosine, 5-methylcytosine, 2,6-diaminopurine, or any other non-naturally occurring base that can be complementary or non-complementary to target RNA or a non-nucleosidic base such as phenyl, naphthyl, 3-nitropyrrole, 5-nitroindole, nebularine, pyridone, pyridinone, or any other non-naturally occurring universal base that can be complementary to target RNA.

The chemically-modified nucleotide or non-nucleotide of Formula II can be present in one or both oligonucleotide strands of the siNA duplex, for example in the sense strand, the antisense strand, or both strands. The siNA molecules of the invention can comprise one or more chemically-modified nucleotide or non-nucleotide of Formula II at the 3'-end, the 5'-end, or both of the 3' and 5'-ends of the sense strand, the antisense strand, or both strands. For example, an exemplary siNA molecule of the invention can comprise about 1 to about 5 or more (e.g., about 1, 2, 3, 4, 5, or more) chemically-modified nucleotides or non-nucleotides of Formula II at the 5'-end of the sense strand, the antisense strand, or both strands. In anther non-limiting example, an exemplary siNA molecule of the invention can comprise about 1 to about 5 or more (e.g., about 1, 2, 3, 4, 5, or more) chemically-modified nucleotides or non-nucleotides of Formula II at the 3'-end of the sense strand, the antisense strand, or both strands.

In one embodiment, the invention features a chemically-modified short interfering nucleic acid (siNA) molecule capable of mediating RNA interference (RNAi) inside a cell or reconstituted *in vitro* system, wherein the chemical modification comprises one or more (e.g., about 1, 2, 3, 4, 5, 6, 7, 8, 9, 10, or more) nucleotides or non-nucleotides having Formula III:

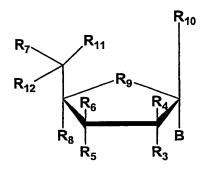
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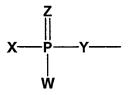
wherein each R3, R4, R5, R6, R7, R8, R10, R11 and R12 is independently H, OH, alkyl, substituted alkyl, alkaryl or aralkyl, F, Cl, Br, CN, CF3, OCF3, OCN, O-alkyl, S-alkyl, N-alkyl, O-alkyl, S-alkyl, N-alkyl, O-alkyl-OH, O-alkyl-OH, O-alkyl-OH, S-alkyl-SH, S-alkyl-OH, S-alkyl-SH, alkyl-S-alkyl, alkyl-O-alkyl, ONO2, NO2, N3, NH2, aminoalkyl, aminoacid, aminoacyl, ONH2, O-aminoalkyl, O-aminoacid, O-aminoacyl, heterocycloalkyl, heterocycloalkaryl, aminoalkylamino, polyalklylamino, substituted silyl, or group having Formula I; R9 is O, S, CH2, S=O, CHF, or CF2, and B is a nucleosidic base such as adenine, guanine, uracil, cytosine, thymine, 2-aminoadenosine, 5-methylcytosine, 2,6-diaminopurine, or any other non-naturally occurring base that can be employed to be complementary or non-complementary to target RNA or a non-nucleosidic base such as phenyl, naphthyl, 3-nitropyrrole, 5-nitroindole, nebularine, pyridone, pyridinone, or any other non-naturally occurring universal base that can be complementary or non-complementary to target RNA.

The chemically-modified nucleotide or non-nucleotide of Formula III can be present in one or both oligonucleotide strands of the siNA duplex, for example, in the sense strand, the antisense strand, or both strands. The siNA molecules of the invention can comprise one or more chemically-modified nucleotide or non-nucleotide of Formula III at the 3'-end, the 5'-end, or both of the 3' and 5'-ends of the sense strand, the antisense strand, or both strands. For example, an exemplary siNA molecule of the invention can comprise about 1 to about 5 or more (e.g., about 1, 2, 3, 4, 5, or more) chemically-

modified nucleotide(s) or non-nucleotide(s) of Formula III at the 5'-end of the sense strand, the antisense strand, or both strands. In anther non-limiting example, an exemplary siNA molecule of the invention can comprise about 1 to about 5 or more (e.g., about 1, 2, 3, 4, 5, or more) chemically-modified nucleotide or non-nucleotide of Formula III at the 3'-end of the sense strand, the antisense strand, or both strands.

In another embodiment, a siNA molecule of the invention comprises a nucleotide having Formula II or III, wherein the nucleotide having Formula II or III is in an inverted configuration. For example, the nucleotide having Formula II or III is connected to the siNA construct in a 3'-3', 3'-2', 2'-3', or 5'-5' configuration, such as at the 3'-end, the 5'-end, or both of the 3' and 5'-ends of one or both siNA strands.

In one embodiment, the invention features a chemically-modified short interfering nucleic acid (siNA) molecule capable of mediating RNA interference (RNAi) inside a cell or reconstituted *in vitro* system, wherein the chemical modification comprises a 5'-terminal phosphate group having Formula IV:



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wherein each X and Y is independently O, S, N, alkyl, substituted alkyl, or alkylhalo; wherein each Z and W is independently O, S, N, alkyl, substituted alkyl, O-alkyl, S-alkyl, alkaryl, aralkyl, or alkylhalo; and wherein W, X, Y and Z are not all O.

In one embodiment, the invention features a siNA molecule having a 5'-terminal phosphate group having Formula IV on the target-complementary strand, for example, a strand complementary to a target RNA, wherein the siNA molecule comprises an all RNA siNA molecule. In another embodiment, the invention features a siNA molecule having a 5'-terminal phosphate group having Formula IV on the target-complementary strand wherein the siNA molecule also comprises about 1 to about 3 (e.g., about 1, 2, or 3) nucleotide 3'-terminal nucleotide overhangs having about 1 to about 4 (e.g., about 1, 2, 3, or 4) deoxyribonucleotides on the 3'-end of one or both strands. In another embodiment, a 5'-terminal phosphate group having Formula IV is present on the target-complementary

strand of a siNA molecule of the invention, for example a siNA molecule having chemical modifications having any of Formulae I-VII.

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In one embodiment, the invention features a chemically-modified short interfering nucleic acid (siNA) molecule capable of mediating RNA interference (RNAi) inside a cell or reconstituted in vitro system, wherein the chemical modification comprises one or more phosphorothioate internucleotide linkages. For example, in a non-limiting example, the invention features a chemically-modified short interfering nucleic acid (siNA) having about 1, 2, 3, 4, 5, 6, 7, 8 or more phosphorothioate internucleotide linkages in one siNA strand. In yet another embodiment, the invention features a chemically-modified short interfering nucleic acid (siNA) individually having about 1, 2, 3, 4, 5, 6, 7, 8 or more phosphorothioate internucleotide linkages in both siNA strands. The phosphorothioate internucleotide linkages can be present in one or both oligonucleotide strands of the siNA duplex, for example in the sense strand, the antisense strand, or both strands. The siNA molecules of the invention can comprise one or more phosphorothioate internucleotide linkages at the 3'-end, the 5'-end, or both of the 3'- and 5'-ends of the sense strand, the antisense strand, or both strands. For example, an exemplary siNA molecule of the invention can comprise about 1 to about 5 or more (e.g., about 1, 2, 3, 4, 5, or more) consecutive phosphorothioate internucleotide linkages at the 5'-end of the sense strand, the antisense strand, or both strands. In another non-limiting example, an exemplary siNA molecule of the invention can comprise one or more (e.g., about 1, 2, 3, 4, 5, 6, 7, 8, 9, 10, or more) pyrimidine phosphorothioate internucleotide linkages in the sense strand, the antisense strand, or both strands. In yet another non-limiting example, an exemplary siNA molecule of the invention can comprise one or more (e.g., about 1, 2, 3, 4, 5, 6, 7, 8, 9, 10, or more) purine phosphorothioate internucleotide linkages in the sense strand, the antisense strand, or both strands.

In one embodiment, the invention features a siNA molecule, wherein the sense strand comprises one or more, for example, about 1, 2, 3, 4, 5, 6, 7, 8, 9, 10, or more phosphorothicate internucleotide linkages, and/or one or more (e.g., about 1, 2, 3, 4, 5, 6, 7, 8, 9, 10 or more) 2'-deoxy, 2'-O-methyl, 2'-deoxy-2'-fluoro, and/or about one or more (e.g., about 1, 2, 3, 4, 5, 6, 7, 8, 9, 10 or more) universal base modified nucleotides, and optionally a terminal cap molecule at the 3'-end, the 5'-end, or both of the 3'- and 5'-ends of the sense strand; and wherein the antisense strand comprises about 1 to about 10 or

more, specifically about 1, 2, 3, 4, 5, 6, 7, 8, 9, 10, or more phosphorothioate internucleotide linkages, and/or one or more (e.g., about 1, 2, 3, 4, 5, 6, 7, 8, 9, 10 or more) 2'-deoxy, 2'-O-methyl, 2'-deoxy-2'-fluoro, and/or one or more (e.g., about 1, 2, 3, 4, 5, 6, 7, 8, 9, 10 or more) universal base modified nucleotides, and optionally a terminal cap molecule at the 3'-end, the 5'-end, or both of the 3'- and 5'-ends of the antisense strand. In another embodiment, one or more, for example about 1, 2, 3, 4, 5, 6, 7, 8, 9, 10, or more, pyrimidine nucleotides of the sense and/or antisense siNA strand are chemically-modified with 2'-deoxy, 2'-O-methyl and/or 2'-deoxy-2'-fluoro nucleotides, with or without one or more, for example about 1, 2, 3, 4, 5, 6, 7, 8, 9, 10, or more, phosphorothioate internucleotide linkages and/or a terminal cap molecule at the 3'-end, the 5'-end, or both of the 3'- and 5'-ends, being present in the same or different strand.

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In another embodiment, the invention features a siNA molecule, wherein the sense strand comprises about 1 to about 5, specifically about 1, 2, 3, 4, or 5 phosphorothioate internucleotide linkages, and/or one or more (e.g., about 1, 2, 3, 4, 5, or more) 2'-deoxy, 2'-O-methyl, 2'-deoxy-2'-fluoro, and/or one or more (e.g., about 1, 2, 3, 4, 5, or more) universal base modified nucleotides, and optionally a terminal cap molecule at the 3-end, the 5'-end, or both of the 3'- and 5'-ends of the sense strand; and wherein the antisense strand comprises about 1 to about 5 or more, specifically about 1, 2, 3, 4, 5, or more phosphorothioate internucleotide linkages, and/or one or more (e.g., about 1, 2, 3, 4, 5, 6, 7, 8, 9, 10 or more) 2'-deoxy, 2'-O-methyl, 2'-deoxy-2'-fluoro, and/or one or more (e.g., about 1, 2, 3, 4, 5, 6, 7, 8, 9, 10 or more) universal base modified nucleotides, and optionally a terminal cap molecule at the 3'-end, the 5'-end, or both of the 3'- and 5'-ends of the antisense strand. In another embodiment, one or more, for example about 1, 2, 3, 4, 5, 6, 7, 8, 9, 10, or more, pyrimidine nucleotides of the sense and/or antisense siNA strand are chemically-modified with 2'-deoxy, 2'-O-methyl and/or 2'-deoxy-2'-fluoro nucleotides, with or without about 1 to about 5 or more, for example about 1, 2, 3, 4, 5, or more phosphorothioate internucleotide linkages and/or a terminal cap molecule at the 3'end, the 5'-end, or both of the 3'- and 5'-ends, being present in the same or different strand.

In one embodiment, the invention features a siNA molecule, wherein the antisense strand comprises one or more, for example, about 1, 2, 3, 4, 5, 6, 7, 8, 9, 10, or more phosphorothioate internucleotide linkages, and/or about one or more (e.g., about 1, 2, 3,

4, 5, 6, 7, 8, 9, 10 or more) 2'-deoxy, 2'-O-methyl, 2'-deoxy-2'-fluoro, and/or one or more (e.g., about 1, 2, 3, 4, 5, 6, 7, 8, 9, 10 or more) universal base modified nucleotides, and optionally a terminal cap molecule at the 3'-end, the 5'-end, or both of the 3'- and 5'-ends of the sense strand; and wherein the antisense strand comprises about 1 to about 10 or more, specifically about 1, 2, 3, 4, 5, 6, 7, 8, 9, 10 or more phosphorothioate internucleotide linkages, and/or one or more (e.g., about 1, 2, 3, 4, 5, 6, 7, 8, 9, 10 or more) 2'-deoxy, 2'-O-methyl, 2'-deoxy-2'-fluoro, and/or one or more (e.g., about 1, 2, 3, 4, 5, 6, 7, 8, 9, 10 or more) universal base modified nucleotides, and optionally a terminal cap molecule at the 3'-end, the 5'-end, or both of the 3'- and 5'-ends of the antisense strand. In another embodiment, one or more, for example about 1, 2, 3, 4, 5, 6, 7, 8, 9, 10 or more pyrimidine nucleotides of the sense and/or antisense siNA strand are chemically-modified with 2'-deoxy, 2'-O-methyl and/or 2'-deoxy-2'-fluoro nucleotides, with or without one or more, for example, about 1, 2, 3, 4, 5, 6, 7, 8, 9, 10 or more phosphorothioate internucleotide linkages and/or a terminal cap molecule at the 3'-end, the 5'-end, or both of the 3' and 5'-ends, being present in the same or different strand.

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In another embodiment, the invention features a siNA molecule, wherein the antisense strand comprises about 1 to about 5 or more, specifically about 1, 2, 3, 4, 5 or more phosphorothioate internucleotide linkages, and/or one or more (e.g., about 1, 2, 3, 4, 5, 6, 7, 8, 9, 10 or more) 2'-deoxy, 2'-O-methyl, 2'-deoxy-2'-fluoro, and/or one or more (e.g., about 1, 2, 3, 4, 5, 6, 7, 8, 9, 10 or more) universal base modified nucleotides, and optionally a terminal cap molecule at the 3'-end, the 5'-end, or both of the 3'- and 5'-ends of the sense strand; and wherein the antisense strand comprises about 1 to about 5 or more, specifically about 1, 2, 3, 4, 5 or more phosphorothioate internucleotide linkages, and/or one or more (e.g., about 1, 2, 3, 4, 5, 6, 7, 8, 9, 10 or more) 2'-deoxy, 2'-O-methyl, 2'-deoxy-2'-fluoro, and/or one or more (e.g., about 1, 2, 3, 4, 5, 6, 7, 8, 9, 10 or more) universal base modified nucleotides, and optionally a terminal cap molecule at the 3'-end, the 5'-end, or both of the 3'- and 5'-ends of the antisense strand. In another embodiment, one or more, for example about 1, 2, 3, 4, 5, 6, 7, 8, 9, 10 or more pyrimidine nucleotides of the sense and/or antisense siNA strand are chemically-modified with 2'-deoxy, 2'-Omethyl and/or 2'-deoxy-2'-fluoro nucleotides, with or without about 1 to about 5, for example about 1, 2, 3, 4, 5 or more phosphorothicate internucleotide linkages and/or a

terminal cap molecule at the 3'-end, the 5'-end, or both of the 3'- and 5'-ends, being present in the same or different strand.

In one embodiment, the invention features a chemically-modified short interfering nucleic acid (siNA) molecule having about 1 to about 5, specifically about 1, 2, 3, 4, 5 or more phosphorothicate internucleotide linkages in each strand of the siNA molecule.

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In another embodiment, the invention features a siNA molecule comprising 2'-5' internucleotide linkages. The 2'-5' internucleotide linkage(s) can be at the 3'-end, the 5'-end, or both of the 3'- and 5'-ends of one or both siNA sequence strands. In addition, the 2'-5' internucleotide linkage(s) can be present at various other positions within one or both siNA sequence strands, for example, about 1, 2, 3, 4, 5, 6, 7, 8, 9, 10, or more including every internucleotide linkage of a pyrimidine nucleotide in one or both strands of the siNA molecule can comprise a 2'-5' internucleotide linkage, or about 1, 2, 3, 4, 5, 6, 7, 8, 9, 10, or more including every internucleotide linkage of a purine nucleotide in one or both strands of the siNA molecule can comprise a 2'-5' internucleotide linkage.

In another embodiment, a chemically-modified siNA molecule of the invention comprises a duplex having two strands, one or both of which can be chemically-modified, wherein each strand is about 18 to about 27 (e.g., about 18, 19, 20, 21, 22, 23, 24, 25, 26, or 27) nucleotides in length, wherein the duplex has about 18 to about 23 (e.g., about 18, 19, 20, 21, 22, or 23) base pairs, and wherein the chemical modification comprises a structure having any of Formulae I-VII. For example, an exemplary chemically-modified siNA molecule of the invention comprises a duplex having two strands, one or both of which can be chemically-modified with a chemical modification having any of Formulae I-VII or any combination thereof, wherein each strand consists of about 21 nucleotides, each having a 2-nucleotide 3'-terminal nucleotide overhang, and wherein the duplex has about 19 base pairs. In another embodiment, a siNA molecule of the invention comprises a single stranded hairpin structure, wherein the siNA is about 36 to about 70 (e.g., about 36, 40, 45, 50, 55, 60, 65, or 70) nucleotides in length having about 18 to about 23 (e.g., about 18, 19, 20, 21, 22, or 23) base pairs, and wherein the siNA can include a chemical modification comprising a structure having any of Formulae I-VII or any combination thereof. For example, an exemplary chemically-modified siNA molecule of the invention comprises a linear oligonucleotide having about 42 to about 50 (e.g., about 42, 43, 44, 45,

46, 47, 48, 49, or 50) nucleotides that is chemically-modified with a chemical modification having any of Formulae I-VII or any combination thereof, wherein the linear oligonucleotide forms a hairpin structure having about 19 base pairs and a 2-nucleotide 3'-terminal nucleotide overhang. In another embodiment, a linear hairpin siNA molecule of the invention contains a stem loop motif, wherein the loop portion of the siNA molecule is biodegradable. For example, a linear hairpin siNA molecule of the invention is designed such that degradation of the loop portion of the siNA molecule in vivo can generate a double-stranded siNA molecule with 3'-terminal overhangs, such as 3'-terminal nucleotide overhangs comprising about 2 nucleotides.

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In another embodiment, a siNA molecule of the invention comprises a circular nucleic acid molecule, wherein the siNA is about 38 to about 70 (e.g., about 38, 40, 45, 50, 55, 60, 65, or 70) nucleotides in length having about 18 to about 23 (e.g., about 18, 19, 20, 21, 22, or 23) base pairs, and wherein the siNA can include a chemical modification, which comprises a structure having any of Formulae I-VII or any combination thereof. For example, an exemplary chemically-modified siNA molecule of the invention comprises a circular oligonucleotide having about 42 to about 50 (e.g., about 42, 43, 44, 45, 46, 47, 48, 49, or 50) nucleotides that is chemically-modified with a chemical modification having any of Formulae I-VII or any combination thereof, wherein the circular oligonucleotide forms a dumbbell shaped structure having about 19 base pairs and 2 loops.

In another embodiment, a circular siNA molecule of the invention contains two loop motifs, wherein one or both loop portions of the siNA molecule is biodegradable. For example, a circular siNA molecule of the invention is designed such that degradation of the loop portions of the siNA molecule *in vivo* can generate a double-stranded siNA molecule with 3'-terminal overhangs, such as 3'-terminal nucleotide overhangs comprising about 2 nucleotides.

In one embodiment, a siNA molecule of the invention comprises at least one (e.g., about 1, 2, 3, 4, 5, 6, 7, 8, 9, 10, or more) abasic moiety, for example a compound having Formula V:

$$R_{7}$$
 R_{12}
 R_{6}
 R_{9}
 R_{4}
 R_{13}

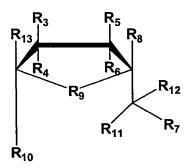
wherein each R3, R4, R5, R6, R7, R8, R10, R11, R12, and R13 is independently H, OH, alkyl, substituted alkyl, alkaryl or aralkyl, F, Cl, Br, CN, CF3, OCF3, OCN, O-alkyl, S-alkyl, N-alkyl, O-alkyl, S-alkyl, N-alkyl, S-alkyl, S-alkyl-OH, O-alkyl-OH, O-alkyl-OH, S-alkyl-SH, alkyl-S-alkyl, alkyl-O-alkyl, ONO2, NO2, N3, NH2, aminoalkyl, aminoacid, aminoacyl, ONH2, O-aminoalkyl, O-aminoacid, O-aminoacyl, heterocycloalkyl, heterocycloalkaryl, aminoalkylamino, polyalklylamino, substituted silyl, or group having Formula I; R9 is O, S, CH2, S=O, CHF, or CF2.

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In one embodiment, a siNA molecule of the invention comprises at least one (e.g., about 1, 2, 3, 4, 5, 6, 7, 8, 9, 10, or more) inverted abasic moiety, for example a compound having Formula VI:



wherein each R3, R4, R5, R6, R7, R8, R10, R11, R12, and R13 is independently H, OH, alkyl, substituted alkyl, alkaryl or aralkyl, F, Cl, Br, CN, CF3, OCF3, OCN, O-alkyl, S-alkyl, N-alkyl, O-alkyl, S-alkyl, N-alkyl-OH, O-alkyl-OH, O-alkyl-OH, O-alkyl-SH, S-alkyl-OH, S-alkyl-SH, alkyl-S-alkyl, alkyl-O-alkyl, ONO2, NO2, N3, NH2, aminoalkyl, aminoacid, aminoacyl, ONH2, O-aminoalkyl, O-aminoacid, O-aminoacyl, heterocycloalkyl, heterocycloalkaryl, aminoalkylamino, polyalklylamino, substituted silyl, or group having Formula I; R9 is O, S, CH2, S=O, CHF, or CF2, and

either R2, R3, R8 or R13 serve as points of attachment to the siNA molecule of the invention.

In another embodiment, a siNA molecule of the invention comprises at least one (e.g., about 1, 2, 3, 4, 5, 6, 7, 8, 9, 10, or more) substituted polyalkyl moieties, for example a compound having Formula VII:

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$$R_1$$
 n
 R_2

wherein each n is independently an integer from 1 to 12, each R1, R2 and R3 is independently H, OH, alkyl, substituted alkyl, alkaryl or aralkyl, F, Cl, Br, CN, CF3, OCF3, OCN, O-alkyl, S-alkyl, N-alkyl, O-alkenyl, S-alkenyl, N-alkenyl, SO-alkyl, alkyl-OSH, alkyl-OH, O-alkyl-OH, O-alkyl-SH, S-alkyl-OH, S-alkyl-SH, alkyl-S-alkyl, alkyl-O-alkyl, ONO2, NO2, N3, NH2, aminoalkyl, aminoacid, aminoacyl, ONH2, O-aminoalkyl, O-aminoacid, O-aminoacyl, heterocycloalkyl, heterocycloalkaryl, aminoalkylamino, polyalklylamino, substituted silyl, or a group having Formula I, and R1, R2 or R3 serves as points of attachment to the siNA molecule of the invention.

In another embodiment, the invention features a compound having Formula VII, wherein R1 and R2 are hydroxyl (OH) groups, n = 1, and R3 comprises O and is the point of attachment to the 3'-end, the 5'-end, or both of the 3' and 5'-ends of one or both strands of a double-stranded siNA molecule of the invention or to a single-stranded siNA molecule of the invention. This modification is referred to herein as "glyceryl" (for example modification 6 in Figure 22).

In another embodiment, a moiety having any of Formula V, VI or VII of the invention is at the 3'-end, the 5'-end, or both of the 3' and 5'-ends of a siNA molecule of the invention. For example, a moiety having Formula V, VI or VII can be present at the 3'-end, the 5'-end, or both of the 3' and 5'-ends of the antisense strand, the sense strand, or both antisense and sense strands of the siNA molecule. In addition, a moiety having Formula VII can be present at the 3'-end or the 5'-end of a hairpin siNA molecule as described herein.

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In another embodiment, a siNA molecule of the invention comprises an abasic residue having Formula V or VI, wherein the abasic residue having Formula VI or VI is connected to the siNA construct in a 3'-3', 3'-2', 2'-3', or 5'-5' configuration, such as at the 3'-end, the 5'-end, or both of the 3' and 5'-ends of one or both siNA strands.

In one embodiment, a siNA molecule of the invention comprises one or more (e.g., about 1, 2, 3, 4, 5, 6, 7, 8, 9, 10, or more) locked nucleic acid (LNA) nucleotides, for example at the 5'-end, the 3'-end, both of the 5' and 3'-ends, or any combination thereof, of the siNA molecule.

In another embodiment, a siNA molecule of the invention comprises one or more (e.g., about 1, 2, 3, 4, 5, 6, 7, 8, 9, 10, or more) acyclic nucleotides, for example at the 5'-end, the 3'-end, both of the 5' and 3'-ends, or any combination thereof, of the siNA molecule.

In one embodiment, the invention features a chemically-modified short interfering nucleic acid (siNA) molecule of the invention, wherein the chemically-modified siNA comprises a sense region, where any (e.g., one or more or all) pyrimidine nucleotides present in the sense region are 2'-deoxy-2'-fluoro pyrimidine nucleotides (e.g., wherein all pyrimidine nucleotides are 2'-deoxy-2'-fluoro pyrimidine nucleotides or alternately a plurality of pyrimidine nucleotides are 2'-deoxy-2'-fluoro pyrimidine nucleotides), and where any (e.g., one or more or all) purine nucleotides present in the sense region are 2'-deoxy purine nucleotides (e.g., wherein all purine nucleotides are 2'-deoxy purine nucleotides or alternately a plurality of purine nucleotides are 2'-deoxy purine nucleotides).

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In one embodiment, the invention features a chemically-modified short interfering nucleic acid (siNA) molecule of the invention, wherein the chemically-modified siNA comprises a sense region, where any (e.g., one or more or all) pyrimidine nucleotides present in the sense region are 2'-deoxy-2'-fluoro pyrimidine nucleotides (e.g., wherein all pyrimidine nucleotides are 2'-deoxy-2'-fluoro pyrimidine nucleotides or alternately a plurality of pyrimidine nucleotides are 2'-deoxy-2'-fluoro pyrimidine nucleotides), and where any (e.g., one or more or all) purine nucleotides present in the sense region are 2'-deoxy purine nucleotides (e.g., wherein all purine nucleotides are 2'-deoxy purine nucleotides or alternately a plurality of purine nucleotides are 2'-deoxy purine

nucleotides), wherein any nucleotides comprising a 3'-terminal nucleotide overhang that are present in said sense region are 2'-deoxy nucleotides.

In one embodiment, the invention features a chemically-modified short interfering nucleic acid (siNA) molecule of the invention, wherein the chemically-modified siNA comprises an antisense region, where any (e.g., one or more or all) pyrimidine nucleotides present in the antisense region are 2'-deoxy-2'-fluoro pyrimidine nucleotides (e.g., wherein all pyrimidine nucleotides are 2'-deoxy-2'-fluoro pyrimidine nucleotides or alternately a plurality of pyrimidine nucleotides are 2'-deoxy-2'-fluoro pyrimidine nucleotides), and wherein any (e.g., one or more or all) purine nucleotides present in the antisense region are 2'-O-methyl purine nucleotides (e.g., wherein all purine nucleotides are 2'-O-methyl purine nucleotides or alternately a plurality of purine nucleotides are 2'-O-methyl purine nucleotides).

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In one embodiment, the invention features a chemically-modified short interfering nucleic acid (siNA) molecule of the invention, wherein the chemically-modified siNA comprises an antisense region, where any (e.g., one or more or all) pyrimidine nucleotides present in the antisense region are 2'-deoxy-2'-fluoro pyrimidine nucleotides (e.g., wherein all pyrimidine nucleotides are 2'-deoxy-2'-fluoro pyrimidine nucleotides or alternately a plurality of pyrimidine nucleotides are 2'-deoxy-2'-fluoro pyrimidine nucleotides), and wherein any (e.g., one or more or all) purine nucleotides present in the antisense region are 2'-O-methyl purine nucleotides (e.g., wherein all purine nucleotides are 2'-O-methyl purine nucleotides or alternately a plurality of purine nucleotides are 2'-O-methyl purine nucleotides or alternately a plurality of purine nucleotides are 2'-O-methyl purine nucleotides), wherein any nucleotides comprising a 3'-terminal nucleotide overhang that are present in said antisense region are 2'-deoxy nucleotides.

In one embodiment, the invention features a chemically-modified short interfering nucleic acid (siNA) molecule of the invention, wherein the chemically-modified siNA comprises an antisense region, where any (e.g., one or more or all) pyrimidine nucleotides present in the antisense region are 2'-deoxy-2'-fluoro pyrimidine nucleotides (e.g., wherein all pyrimidine nucleotides are 2'-deoxy-2'-fluoro pyrimidine nucleotides or alternately a plurality of pyrimidine nucleotides are 2'-deoxy-2'-fluoro pyrimidine nucleotides), and where any (e.g., one or more or all) purine nucleotides present in the antisense region are 2'-deoxy purine nucleotides (e.g., wherein all purine nucleotides are

2'-deoxy purine nucleotides or alternately a plurality of purine nucleotides are 2'-deoxy purine nucleotides).

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In one embodiment, the invention features a chemically-modified short interfering nucleic acid (siNA) molecule of the invention capable of mediating RNA interference (RNAi) inside a cell or reconstituted in vitro system, wherein the chemically-modified siNA comprises a sense region and an antisense region. The sense region comprises one 2'-deoxy-2'-fluoro pyrimidine nucleotides (e.g., wherein all pyrimidine nucleotides are 2'-deoxy-2'-fluoro pyrimidine nucleotides or alternately a plurality of pyrimidine nucleotides are 2'-deoxy-2'-fluoro pyrimidine nucleotides), and one or more 2'-deoxy purine nucleotides (e.g., wherein all purine nucleotides are 2'-deoxy purine nucleotides or alternately a plurality of purine nucleotides are 2'-deoxy purine nucleotides). Inverted deoxy abasic modifications can be optionally present at the 3'end, the 5'-end, or both of the 3' and 5'-ends of the sense region. The sense region optionally further comprises a 3'-terminal overhang having about 1 to about 4 (e.g., about 1, 2, 3, or 4) 2'-deoxyribonucleotides. The antisense region comprises one or more 2'deoxy-2'-fluoro pyrimidine nucleotides (e.g., wherein all pyrimidine nucleotides are 2'deoxy-2'-fluoro pyrimidine nucleotides or alternately a plurality of pyrimidine nucleotides are 2'-deoxy-2'-fluoro pyrimidine nucleotides), and one or more 2'-O-methyl purine nucleotides (e.g., wherein all purine nucleotides are 2'-O-methyl purine nucleotides or alternately a plurality of purine nucleotides are 2'-O-methyl purine nucleotides). A terminal cap modification, such as any modification described herein or shown in Figure 22, is optionally present at the 3'-end, the 5'-end, or both of the 3' and 5'-ends of the antisense sequence. The antisense region optionally further comprises a 3'-terminal nucleotide overhang having about 1 to about 4 (e.g., about 1, 2, 3, or 4) 2'deoxynucleotides, wherein the overhang nucleotides can further comprise one or more (e.g., 1, 2, 3, or 4) phosphorothioate internucleotide linkages. Non-limiting examples of these chemically-modified siNAs are shown in Figures 18 and 19 and Table IV herein.

In one embodiment, the invention features a chemically-modified short interfering nucleic acid (siNA) molecule of the invention capable of mediating RNA interference (RNAi) inside a cell or reconstituted *in vitro* system, wherein the siNA comprises a sense region and an antisense region, wherein the sense region comprises one or more 2'-deoxy-2'-fluoro pyrimidine nucleotides (e.g., wherein all pyrimidine nucleotides are 2'-deoxy-2'-

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fluoro pyrimidine nucleotides or alternately a plurality of pyrimidine nucleotides are 2'deoxy-2'-fluoro pyrimidine nucleotides), and one or more purine ribonucleotides (e.g., wherein all purine nucleotides are purine ribonucleotides or alternately a plurality of purine nucleotides are purine ribonucleotides) and wherein the antisense region comprises one or more 2'-deoxy-2'-fluoro pyrimidine nucleotides (e.g., wherein all pyrimidine nucleotides are 2'-deoxy-2'-fluoro pyrimidine nucleotides or alternately a plurality of pyrimidine nucleotides are 2'-deoxy-2'-fluoro pyrimidine nucleotides), and one or more 2'-O-methyl purine nucleotides (e.g., wherein all purine nucleotides are 2'-O-methyl purine nucleotides or alternately a plurality of purine nucleotides are 2'-O-methyl purine nucleotides). Inverted deoxy abasic modifications are optionally present at the 3'-end, the 5'-end, or both of the 3' and 5'-ends of the sense region. The sense region optionally further comprises a 3'-terminal overhang having about 1 to about 4 (e.g., about 1, 2, 3, or 4) 2'-deoxyribonucleotides. A terminal cap modification, such as any modification described herein or shown in Figure 22, is optionally present at the 3'-end, the 5'-end, or both of the 3' and 5'-ends of the antisense sequence. The antisense region optionally further comprises a 3'-terminal nucleotide overhang having about 1 to about 4 (e.g., about 1, 2, 3, or 4) 2'-deoxynucleotides, wherein the overhang nucleotides can further comprise one or more (e.g., 1, 2, 3, or 4) phosphorothioate internucleotide linkages. Non-limiting examples of these chemically-modified siNAs are shown in Figures 18 and 19 and Table IV herein.

In one embodiment, the invention features a chemically-modified short interfering nucleic acid (siNA) molecule of the invention capable of mediating RNA interference (RNAi) inside a cell or reconstituted *in vitro* system, wherein the chemically-modified siNA comprises a sense region and an antisense region, wherein the sense region comprises one or 2'-deoxy-2'-fluoro pyrimidine nucleotides (e.g., wherein all pyrimidine nucleotides are 2'-deoxy-2'-fluoro pyrimidine nucleotides or alternately a plurality of pyrimidine nucleotides are 2'-deoxy-2'-fluoro pyrimidine nucleotides), and one or more purine nucleotides selected from the group consisting of 2'-deoxy nucleotides, locked nucleic acid (LNA) nucleotides, 2'-methoxyethyl nucleotides are selected from the group consisting of 2'-deoxy nucleotides, locked nucleic acid (LNA) nucleotides, 2'-methoxyethyl nucleotides are selected from the group consisting of 2'-deoxy nucleotides, locked nucleic acid (LNA) nucleotides, 2'-methoxyethyl nucleotides or alternately methoxyethyl nucleotides, 4'-thionucleotides, and 2'-O-methyl nucleotides or alternately

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a plurality of purine nucleotides are selected from the group consisting of 2'-deoxy nucleotides, locked nucleic acid (LNA) nucleotides, 2'-methoxyethyl nucleotides, 4'thionucleotides, and 2'-O-methyl nucleotides) and wherein the antisense region comprises one or more 2'-deoxy-2'-fluoro pyrimidine nucleotides (e.g., wherein all pyrimidine nucleotides are 2'-deoxy-2'-fluoro pyrimidine nucleotides or alternately a plurality of pyrimidine nucleotides are 2'-deoxy-2'-fluoro pyrimidine nucleotides), and one or more purine nucleotides selected from the group consisting of 2'-deoxy nucleotides, locked nucleic acid (LNA) nucleotides, 2'-methoxyethyl nucleotides, 4'-thionucleotides, and 2'-O-methyl nucleotides (e.g., wherein all purine nucleotides are selected from the group consisting of 2'-deoxy nucleotides, locked nucleic acid (LNA) nucleotides, 2'methoxyethyl nucleotides, 4'-thionucleotides, and 2'-O-methyl nucleotides or alternately a plurality of purine nucleotides are selected from the group consisting of 2'-deoxy nucleotides, locked nucleic acid (LNA) nucleotides, 2'-methoxyethyl nucleotides, 4'thionucleotides, and 2'-O-methyl nucleotides). Inverted deoxy abasic modifications are optionally present at the 3'-end, the 5'-end, or both of the 3' and 5'-ends of the sense region. The sense region optionally further comprises a 3'-terminal overhang having about 1 to about 4 (e.g., about 1, 2, 3, or 4) 2'-deoxyribonucleotides. A terminal cap modification, such as any modification described herein or shown in Figure 22, is optionally present at the 3'-end, the 5'-end, or both of the 3' and 5'-ends of the antisense sequence. The antisense region optionally further comprises a 3'-terminal nucleotide overhang having about 1 to about 4 (e.g., about 1, 2, 3, or 4) 2'-deoxynucleotides, wherein the overhang nucleotides can further comprise one or more (e.g., 1, 2, 3, or 4) phosphorothioate internucleotide linkages.

In another embodiment, any modified nucleotides present in the siNA molecules of the invention, preferably in the antisense strand of the siNA molecules of the invention, but also optionally in the sense and/or both antisense and sense strands, comprise modified nucleotides having properties or characteristics similar to naturally occurring ribonucleotides. For example, the invention features siNA molecules including modified nucleotides having a Northern conformation (e.g., Northern pseudorotation cycle, see for example Saenger, *Principles of Nucleic Acid Structure*, Springer-Verlag ed., 1984). As such, chemically modified nucleotides present in the siNA molecules of the invention, preferably in the antisense strand of the siNA molecules of the invention, but also

optionally in the sense and/or both antisense and sense strands, are resistant to nuclease degradation while at the same time maintaining the capacity to mediate RNAi. Non-limiting examples of nucleotides having a northern configuration include locked nucleic acid (LNA) nucleotides (e.g., 2'-O,4'-C-methylene-(D-ribofuranosyl) nucleotides); 2'-methoxyethoxy (MOE) nucleotides; 2'-methyl-thio-ethyl, 2'-deoxy-2'-fluoro nucleotides, 2'-deoxy-2'-chloro nucleotides, 2'-azido nucleotides, and 2'-O-methyl nucleotides.

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In one embodiment, the invention features a chemically-modified short interfering nucleic acid molecule (siNA) capable of mediating RNA interference (RNAi) inside a cell or reconstituted in vitro system, wherein the chemical modification comprises a conjugate covalently attached to the chemically-modified siNA molecule. In another embodiment, the conjugate is covalently attached to the chemically-modified siNA molecule via a biodegradable linker. In one embodiment, the conjugate molecule is attached at the 3'end of either the sense strand, the antisense strand, or both strands of the chemicallymodified siNA molecule. In another embodiment, the conjugate molecule is attached at the 5'-end of either the sense strand, the antisense strand, or both strands of the chemically-modified siNA molecule. In yet another embodiment, the conjugate molecule is attached both the 3'-end and 5'-end of either the sense strand, the antisense strand, or both strands of the chemically-modified siNA molecule, or any combination thereof. In one embodiment, a conjugate molecule of the invention comprises a molecule that facilitates delivery of a chemically-modified siNA molecule into a biological system, such as a cell. In another embodiment, the conjugate molecule attached to the chemically-modified siNA molecule is a poly ethylene glycol, human serum albumin, or a ligand for a cellular receptor that can mediate cellular uptake. Examples of specific conjugate molecules contemplated by the instant invention that can be attached to chemically-modified siNA molecules are described in Vargeese et al., U.S. Serial No. 10/201,394, incorporated by reference herein. The type of conjugates used and the extent of conjugation of siNA molecules of the invention can be evaluated for improved pharmacokinetic profiles, bioavailability, and/or stability of siNA constructs while at the same time maintaining the ability of the siNA to mediate RNAi activity. As such, one skilled in the art can screen siNA constructs that are modified with various conjugates to determine whether the siNA conjugate complex possesses improved properties while

maintaining the ability to mediate RNAi, for example in animal models as are generally known in the art.

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In one embodiment, the invention features a short interfering nucleic acid (siNA) molecule of the invention, wherein the siNA further comprises a nucleotide, nonnucleotide, or mixed nucleotide/non-nucleotide linker that joins the sense region of the siNA to the antisense region of the siNA. In one embodiment, a nucleotide linker of the invention can be a linker of ≥ 2 nucleotides in length, for example 3, 4, 5, 6, 7, 8, 9, or 10 nucleotides in length. In another embodiment, the nucleotide linker can be a nucleic acid aptamer. By "aptamer" or "nucleic acid aptamer" as used herein is meant a nucleic acid molecule that binds specifically to a target molecule wherein the nucleic acid molecule has sequence that comprises a sequence recognized by the target molecule in its natural setting. Alternately, an aptamer can be a nucleic acid molecule that binds to a target molecule where the target molecule does not naturally bind to a nucleic acid. The target molecule can be any molecule of interest. For example, the aptamer can be used to bind to a ligand-binding domain of a protein, thereby preventing interaction of the naturally occurring ligand with the protein. This is a non-limiting example and those in the art will recognize that other embodiments can be readily generated using techniques generally known in the art. (See, for example, Gold et al., 1995, Annu. Rev. Biochem., 64, 763; Brody and Gold, 2000, J. Biotechnol., 74, 5; Sun, 2000, Curr. Opin. Mol. Ther., 2, 100; Kusser, 2000, J. Biotechnol., 74, 27; Hermann and Patel, 2000, Science, 287, 820; and Jayasena, 1999, Clinical Chemistry, 45, 1628.)

In yet another embodiment, a non-nucleotide linker of the invention comprises abasic nucleotide, polyether, polyamine, polyamide, peptide, carbohydrate, lipid, polyhydrocarbon, or other polymeric compounds (e.g. polyethylene glycols such as those having between 2 and 100 ethylene glycol units). Specific examples include those described by Seela and Kaiser, Nucleic Acids Res. 1990, 18:6353 and Nucleic Acids Res. 1987, 15:3113; Cload and Schepartz, J. Am. Chem. Soc. 1991, 113:6324; Richardson and Schepartz, J. Am. Chem. Soc. 1991, 113:5109; Ma et al., Nucleic Acids Res. 1993, 21:2585 and Biochemistry 1993, 32:1751; Durand et al., Nucleic Acids Res. 1990, 18:6353; McCurdy et al., Nucleosides & Nucleotides 1991, 10:287; Jschke et al., Tetrahedron Lett. 1993, 34:301; Ono et al., Biochemistry 1991, 30:9914; Arnold et al., International Publication No. WO 89/02439; Usman et al., International Publication No.

WO 95/06731; Dudycz et al., International Publication No. WO 95/11910 and Ferentz and Verdine, J. Am. Chem. Soc. 1991, 113:4000, all hereby incorporated by reference herein. A "non-nucleotide" further means any group or compound that can be incorporated into a nucleic acid chain in the place of one or more nucleotide units, including either sugar and/or phosphate substitutions, and allows the remaining bases to exhibit their enzymatic activity. The group or compound can be abasic in that it does not contain a commonly recognized nucleotide base, such as adenosine, guanine, cytosine, uracil or thymine, for example at the C1 position of the sugar.

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In one embodiment, the invention features a short interfering nucleic acid (siNA) molecule capable of mediating RNA interference (RNAi) inside a cell or reconstituted in vitro system, wherein one or both strands of the siNA molecule that are assembled from two separate oligonucleotides do not comprise any ribonucleotides. For example, a siNA molecule can be assembled from a single oligonculeotide where the sense and antisense regions of the siNA comprise separate oligonucleotides not having any ribonucleotides (e.g., nucleotides having a 2'-OH group) present in the oligonucleotides. In another example, a siNA molecule can be assembled from a single oligonculeotide where the sense and antisense regions of the siNA are linked or circularized by a nucleotide or nonnucleotide linker as desrcibed herein, wherein the oligonucleotide does not have any ribonucleotides (e.g., nucleotides having a 2'-OH group) present in the oligonucleotide. Applicant has surprisingly found that the presense of ribonucleotides (e.g., nucleotides having a 2'-hydroxyl group) within the siNA molecule is not required or essential to support RNAi activity. As such, in one embodiment, all positions within the siNA can include chemically modified nucleotides and/or non-nucleotides such as nucleotides and or non-nucleotides having Formula I, II, III, IV, V, VI, or VII or any combination thereof to the extent that the ability of the siNA molecule to support RNAi activity in a cell is maintained.

In one embodiment, a siNA molecule of the invention is a single stranded siNA molecule that mediates RNAi activity in a cell or reconstituted in vitro system, wherein the siNA molecule comprises a single stranded polynucleotide having complementarity to a target nucleic acid sequence. In another embodiment, the single stranded siNA molecule of the invention comprises a 5'-terminal phosphate group. In another embodiment, the single stranded siNA molecule of the invention comprises a 5'-terminal

phosphate group and a 3'-terminal phosphate group (e.g., a 2', 3'-cyclic phosphate). In another embodiment, the single stranded siNA molecule of the invention comprises about 19 to about 29 nucleotides. In yet another embodiment, the single stranded siNA molecule of the invention comprises one or more chemically modified nucleotides or non-nucleotides described herein. For example, all the positions within the siNA molecule can include chemically-modified nucleotides such as nucleotides having any of Formulae I-VII, or any combination thereof to the extent that the ability of the siNA molecule to support RNAi activity in a cell is maintained.

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In one embodiment, a siNA molecule of the invention is a single stranded siNA molecule that mediates RNAi activity in a cell or reconstituted in vitro system, wherein the siNA molecule comprises a single stranded polynucleotide having complementarity to a target nucleic acid sequence, and wherein one or more pyrimidine nucleotides present in the siNA are 2'-deoxy-2'-fluoro pyrimidine nucleotides (e.g., wherein all pyrimidine nucleotides are 2'-deoxy-2'-fluoro pyrimidine nucleotides or alternately a plurality of pyrimidine nucleotides are 2'-deoxy-2'-fluoro pyrimidine nucleotides), and wherein any purine nucleotides present in the antisense region are 2'-O-methyl purine nucleotides (e.g., wherein all purine nucleotides are 2'-O-methyl purine nucleotides or alternately a plurality of purine nucleotides are 2'-O-methyl purine nucleotides), and a terminal cap modification, such as any modification described herein or shown in Figure 22, that is optionally present at the 3'-end, the 5'-end, or both of the 3' and 5'-ends of the antisense sequence, the siNA optionally further comprising about 1 to about 4 (e.g., about 1, 2, 3, or 4) terminal 2'-deoxynucleotides at the 3'-end of the siNA molecule, wherein the terminal nucleotides can further comprise one or more (e.g., 1, 2, 3, or 4) phosphorothioate internucleotide linkages, and wherein the siNA optionally further comprises a terminal phosphate group, such as a 5'-terminal phosphate group.

In one embodiment, a siNA molecule of the invention is a single stranded siNA molecule that mediates RNAi activity in a cell or reconstituted in vitro system, wherein the siNA molecule comprises a single stranded polynucleotide having complementarity to a target nucleic acid sequence, and wherein one or more pyrimidine nucleotides present in the siNA are 2'-deoxy-2'-fluoro pyrimidine nucleotides (e.g., wherein all pyrimidine nucleotides are 2'-deoxy-2'-fluoro pyrimidine nucleotides or alternately a plurality of pyrimidine nucleotides are 2'-deoxy-2'-fluoro pyrimidine nucleotides), and wherein any

purine nucleotides present in the antisense region are 2'-deoxy purine nucleotides (e.g., wherein all purine nucleotides are 2'-deoxy purine nucleotides or alternately a plurality of purine nucleotides are 2'-deoxy purine nucleotides), and a terminal cap modification, such as any modification described herein or shown in Figure 22, that is optionally present at the 3'-end, the 5'-end, or both of the 3' and 5'-ends of the antisense sequence, the siNA optionally further comprising about 1 to about 4 (e.g., about 1, 2, 3, or 4) terminal 2'-deoxynucleotides at the 3'-end of the siNA molecule, wherein the terminal nucleotides can further comprise one or more (e.g., 1, 2, 3, or 4) phosphorothioate internucleotide linkages, and wherein the siNA optionally further comprises a terminal phosphate group, such as a 5'-terminal phosphate group.

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In one embodiment, a siNA molecule of the invention is a single stranded siNA molecule that mediates RNAi activity in a cell or reconstituted in vitro system, wherein the siNA molecule comprises a single stranded polynucleotide having complementarity to a target nucleic acid sequence, and wherein one or more pyrimidine nucleotides present in the siNA are 2'-deoxy-2'-fluoro pyrimidine nucleotides (e.g., wherein all pyrimidine nucleotides are 2'-deoxy-2'-fluoro pyrimidine nucleotides or alternately a plurality of pyrimidine nucleotides are 2'-deoxy-2'-fluoro pyrimidine nucleotides), and wherein any purine nucleotides present in the antisense region are locked nucleic acid (LNA) nucleotides (e.g., wherein all purine nucleotides are LNA nucleotides or alternately a plurality of purine nucleotides are LNA nucleotides), and a terminal cap modification, such as any modification described herein or shown in Figure 22, that is optionally present at the 3'-end, the 5'-end, or both of the 3' and 5'-ends of the antisense sequence, the siNA optionally further comprising about 1 to about 4 (e.g., about 1, 2, 3, or 4) terminal 2'-deoxynucleotides at the 3'-end of the siNA molecule, wherein the terminal nucleotides can further comprise one or more (e.g., 1, 2, 3, or 4) phosphorothioate internucleotide linkages, and wherein the siNA optionally further comprises a terminal phosphate group, such as a 5'-terminal phosphate group.

In one embodiment, a siNA molecule of the invention is a single stranded siNA molecule that mediates RNAi activity in a cell or reconstituted in vitro system, wherein the siNA molecule comprises a single stranded polynucleotide having complementarity to a target nucleic acid sequence, and wherein one or more pyrimidine nucleotides present in the siNA are 2'-deoxy-2'-fluoro pyrimidine nucleotides (e.g., wherein all pyrimidine

nucleotides are 2'-deoxy-2'-fluoro pyrimidine nucleotides or alternately a plurality of pyrimidine nucleotides are 2'-deoxy-2'-fluoro pyrimidine nucleotides), and wherein any purine nucleotides present in the antisense region are 2'-methoxyethyl purine nucleotides (e.g., wherein all purine nucleotides are 2'-methoxyethyl purine nucleotides or alternately a plurality of purine nucleotides are 2'-methoxyethyl purine nucleotides), and a terminal cap modification, such as any modification described herein or shown in **Figure 22**, that is optionally present at the 3'-end, the 5'-end, or both of the 3' and 5'-ends of the antisense sequence, the siNA optionally further comprising about 1 to about 4 (e.g., about 1, 2, 3, or 4) terminal 2'-deoxynucleotides at the 3'-end of the siNA molecule, wherein the terminal nucleotides can further comprise one or more (e.g., 1, 2, 3, or 4) phosphorothioate internucleotide linkages, and wherein the siNA optionally further comprises a terminal phosphate group, such as a 5'-terminal phosphate group.

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In another embodiment, any modified nucleotides present in the single stranded siNA molecules of the invention comprise modified nucleotides having properties or characteristics similar to naturally occurring ribonucleotides. For example, the invention features siNA molecules including modified nucleotides having a Northern conformation (e.g., Northern pseudorotation cycle, see for example Saenger, *Principles of Nucleic Acid Structure*, Springer-Verlag ed., 1984). As such, chemically modified nucleotides present in the single stranded siNA molecules of the invention are preferably resistant to nuclease degradation while at the same time maintaining the capacity to mediate RNAi.

In one embodiment, the invention features a method for modulating the expression of a gene within a cell comprising: (a) synthesizing a siNA molecule of the invention, which can be chemically-modified, wherein one of the siNA strands comprises a sequence complementary to RNA of the gene; and (b) introducing the siNA molecule into a cell under conditions suitable to modulate the expression of the gene in the cell.

In one embodiment, the invention features a method for modulating the expression of a gene within a cell comprising: (a) synthesizing a siNA molecule of the invention, which can be chemically-modified, wherein one of the siNA strands comprises a sequence complementary to RNA of the gene and wherein the sense strand sequence of the siNA comprises a sequence substantially similar to the sequence of the target RNA;

and (b) introducing the siNA molecule into a cell under conditions suitable to modulate the expression of the gene in the cell.

In another embodiment, the invention features a method for modulating the expression of more than one gene within a cell comprising: (a) synthesizing siNA molecules of the invention, which can be chemically-modified, wherein one of the siNA strands comprises a sequence complementary to RNA of the genes; and (b) introducing the siNA molecules into a cell under conditions suitable to modulate the expression of the genes in the cell.

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In another embodiment, the invention features a method for modulating the expression of more than one gene within a cell comprising: (a) synthesizing a siNA molecule of the invention, which can be chemically-modified, wherein one of the siNA strands comprises a sequence complementary to RNA of the gene and wherein the sense strand sequence of the siNA comprises a sequence substantially similar to the sequence of the target RNA; and (b) introducing the siNA molecules into a cell under conditions suitable to modulate the expression of the genes in the cell.

In one embodiment, siNA molecules of the invention are used as reagents in ex vivo applications. For example, siNA reagents are intoduced into tissue or cells that are transplanted into a subject for therapeutic effect. The cells and/or tissue can be derived from an organism or subject that later receives the explant, or can be derived from another organism or subject prior to transplantation. The siNA molecules can be used to modulate the expression of one or more genes in the cells or tissue, such that the cells or tissue obtain a desired phenotype or are able to perform a function when transplanted in vivo. In one embodiment, certain target cells from a patient are extracted. These extracted cells are contacted with siNAs targeteing a specific nucleotide sequence within the cells under conditions suitable for uptake of the siNAs by these cells (e.g. using delivery reagents such as cationic lipids, liposomes and the like or using techniques such as electroporation to facilitate the delivery of siNAs into cells). The cells are then reintroduced back into the same patient or other patients. Non-limiting examples of ex vivo applications include use in organ/tissue transplant, tissue grafting, or treatment of pulmonary disease (e.g., restenosis) or prevent neointimal hyperplasia and atherosclerosis in vein grafts. Such ex vivo applications may also used to treat conditions associated with

coronary and peripheral bypass graft failure, for example, such methods can be used in conjunction with peripheral vascular bypass graft surgery and coronary artery bypass graft surgery. Additional applications include transplants to treat CNS lesions or injury, including use in treatment of neurodegenerative conditions such as Alzheimer's disease, Parkinson's Disease, Epilepsy, Dementia, Huntington's disease, or amyotrophic lateral sclerosis (ALS).

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In one embodiment, the invention features a method of modulating the expression of a gene in a tissue explant comprising: (a) synthesizing a siNA molecule of the invention, which can be chemically-modified, wherein one of the siNA strands comprises a sequence complementary to RNA of the gene; and (b) introducing the siNA molecule into a cell of the tissue explant derived from a particular organism under conditions suitable to modulate the expression of the gene in the tissue explant. In another embodiment, the method further comprises introducing the tissue explant back into the organism the tissue was derived from or into another organism under conditions suitable to modulate the expression of the gene in that organism.

In one embodiment, the invention features a method of modulating the expression of a gene in a tissue explant comprising: (a) synthesizing a siNA molecule of the invention, which can be chemically-modified, wherein one of the siNA strands comprises a sequence complementary to RNA of the gene and wherein the sense strand sequence of the siNA comprises a sequence substantially similar to the sequence of the target RNA; and (b) introducing the siNA molecule into a cell of the tissue explant derived from a particular organism under conditions suitable to modulate the expression of the gene in the tissue explant. In another embodiment, the method further comprises introducing the tissue explant back into the organism the tissue was derived from or into another organism under conditions suitable to modulate the expression of the gene in that organism.

In another embodiment, the invention features a method of modulating the expression of more than one gene in a tissue explant comprising: (a) synthesizing siNA molecules of the invention, which can be chemically-modified, wherein one of the siNA strands comprises a sequence complementary to RNA of the genes; and (b) introducing the siNA molecules into a cell of the tissue explant derived from a particular organism

under conditions suitable to modulate the expression of the genes in the tissue explant. In another embodiment, the method further comprises introducing the tissue explant back into the organism the tissue was derived from or into another organism under conditions suitable to modulate the expression of the genes in that organism.

In one embodiment, the invention features a method of modulating the expression of a gene in an organism comprising: (a) synthesizing a siNA molecule of the invention, which can be chemically-modified, wherein one of the siNA strands comprises a sequence complementary to RNA of the gene; and (b) introducing the siNA molecule into the organism under conditions suitable to modulate the expression of the gene in the organism.

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In another embodiment, the invention features a method of modulating the expression of more than one gene in an organism comprising: (a) synthesizing siNA molecules of the invention, which can be chemically-modified, wherein one of the siNA strands comprises a sequence complementary to RNA of the genes; and (b) introducing the siNA molecules into the organism under conditions suitable to modulate the expression of the genes in the organism.

In one embodiment, the invention features a method for modulating the expression of a gene within a cell comprising: (a) synthesizing a siNA molecule of the invention, which can be chemically-modified, wherein the siNA comprises a single stranded sequence having complementarity to RNA of the gene; and (b) introducing the siNA molecule into a cell under conditions suitable to modulate the expression of the gene in the cell.

In another embodiment, the invention features a method for modulating the expression of more than one gene within a cell comprising: (a) synthesizing siNA molecules of the invention, which can be chemically-modified, wherein the siNA comprises a single stranded sequence having complementarity to RNA of the gene; and (b) contacting the siNA molecule with a cell in vitro or in vivo under conditions suitable to modulate the expression of the genes in the cell.

In one embodiment, the invention features a method of modulating the expression of a gene in a tissue explant comprising: (a) synthesizing a siNA molecule of the

invention, which can be chemically-modified, wherein the siNA comprises a single stranded sequence having complementarity to RNA of the gene; and (b) contacting the siNA molecule with a cell of the tissue explant derived from a particular organism under conditions suitable to modulate the expression of the gene in the tissue explant. In another embodiment, the method further comprises introducing the tissue explant back into the organism the tissue was derived from or into another organism under conditions suitable to modulate the expression of the gene in that organism.

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In another embodiment, the invention features a method of modulating the expression of more than one gene in a tissue explant comprising: (a) synthesizing siNA molecules of the invention, which can be chemically-modified, wherein the siNA comprises a single stranded sequence having complementarity to RNA of the gene; and (b) introducing the siNA molecules into a cell of the tissue explant derived from a particular organism under conditions suitable to modulate the expression of the genes in the tissue explant. In another embodiment, the method further comprises introducing the tissue explant back into the organism the tissue was derived from or into another organism under conditions suitable to modulate the expression of the genes in that organism.

In one embodiment, the invention features a method of modulating the expression of a gene in an organism comprising: (a) synthesizing a siNA molecule of the invention, which can be chemically-modified, wherein the siNA comprises a single stranded sequence having complementarity to RNA of the gene; and (b) introducing the siNA molecule into the organism under conditions suitable to modulate the expression of the gene in the organism.

In another embodiment, the invention features a method of modulating the expression of more than one gene in an organism comprising: (a) synthesizing siNA molecules of the invention, which can be chemically-modified, wherein the siNA comprises a single stranded sequence having complementarity to RNA of the gene; and (b) introducing the siNA molecules into the organism under conditions suitable to modulate the expression of the genes in the organism.

In one embodiment, the invention features a method of modulating the expression of a gene in an organism comprising contacting the organism with a siNA molecule of the

invention under conditions suitable to modulate the expression of the gene in the organism.

In another embodiment, the invention features a method of modulating the expression of more than one gene in an organism comprising contacting the organism with one or more siNA molecules of the invention under conditions suitable to modulate the expression of the genes in the organism.

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The siNA molecules of the invention can be designed to inhibit target gene expression through RNAi targeting of a variety of RNA molecules. In one embodiment, the siNA molecules of the invention are used to target various RNAs corresponding to a target gene. Non-limiting examples of such RNAs include messenger RNA (mRNA), alternate RNA splice variants of target gene(s), post-transcriptionally modified RNA of target gene(s), pre-mRNA of target gene(s), and/or RNA templates. If alternate splicing produces a family of transcripts that are distinguished by usage of appropriate exons, the instant invention can be used to inhibit gene expression through the appropriate exons to specifically inhibit or to distinguish among the functions of gene family members. For example, a protein that contains an alternatively spliced transmembrane domain can be expressed in both membrane bound and secreted forms. Use of the invention to target the exon containing the transmembrane domain can be used to determine the functional consequences of pharmaceutical targeting of membrane bound as opposed to the secreted form of the protein. Non-limiting examples of applications of the invention relating to targeting these RNA molecules include therapeutic pharmaceutical applications, pharmaceutical discovery applications, molecular diagnostic and gene function applications, and gene mapping, for example using single nucleotide polymorphism mapping with siNA molecules of the invention. Such applications can be implemented using known gene sequences or from partial sequences available from an expressed sequence tag (EST).

In another embodiment, the siNA molecules of the invention are used to target conserved sequences corresponding to a gene family or gene families. As such, siNA molecules targeting multiple gene targets can provide increased therapeutic effect. In addition, siNA can be used to characterize pathways of gene function in a variety of applications. For example, the present invention can be used to inhibit the activity of

target gene(s) in a pathway to determine the function of uncharacterized gene(s) in gene function analysis, mRNA function analysis, or translational analysis. The invention can be used to determine potential target gene pathways involved in various diseases and conditions toward pharmaceutical development. The invention can be used to understand pathways of gene expression involved in, for example, in development, such as prenatal development and postnatal development, and/or the progression and/or maintenance of cancer, infectious disease, autoimmunity, inflammation, endocrine disorders, renal disease, pulmonary disease, cardiovascular disease, birth defects, ageing, any other disease or condition related to gene expression.

In one embodiment, the invention features a method comprising: (a) generating a library of siNA constructs having a predetermined complexity; and (b) assaying the siNA constructs of (a) above, under conditions suitable to determine RNAi target sites within the target RNA sequence. In another embodiment, the siNA molecules of (a) have strands of a fixed length, for example, about 23 nucleotides in length. In yet another embodiment, the siNA molecules of (a) are of differing length, for example having strands of about 19 to about 25 (e.g., about 19, 20, 21, 22, 23, 24, or 25) nucleotides in length. In one embodiment, the assay can comprise a reconstituted in vitro siNA assay as described herein. In another embodiment, the assay can comprise a cell culture system in which target RNA is expressed. In another embodiment, fragments of target RNA are analyzed for detectable levels of cleavage, for example by gel electrophoresis, northern blot analysis, or RNAse protection assays, to determine the most suitable target site(s) within the target RNA sequence. The target RNA sequence can be obtained as is known in the art, for example, by cloning and/or transcription for in vitro systems, and by cellular expression in in vivo systems.

In one embodiment, the invention features a method comprising: (a) generating a randomized library of siNA constructs having a predetermined complexity, such as of 4^N, where N represents the number of base paired nucleotides in each of the siNA construct strands (eg. for a siNA construct having 21 nucleotide sense and antisense strands with 19 base pairs, the complexity would be 4¹⁹); and (b) assaying the siNA constructs of (a) above, under conditions suitable to determine RNAi target sites within the target RNA sequence. In another embodiment, the siNA molecules of (a) have strands of a fixed length, for example about 23 nucleotides in length. In yet another embodiment, the siNA

molecules of (a) are of differing length, for example having strands of about 19 to about 25 (e.g., about 19, 20, 21, 22, 23, 24, or 25) nucleotides in length. In one embodiment, the assay can comprise a reconstituted in vitro siNA assay as described in Example 7 herein. In another embodiment, the assay can comprise a cell culture system in which target RNA is expressed. In another embodiment, fragments of target RNA are analyzed for detectable levels of cleavage, for example by gel electrophoresis, northern blot analysis, or RNAse protection assays, to determine the most suitable target site(s) within the target RNA sequence. In another embodiment, the target RNA sequence can be obtained as is known in the art, for example, by cloning and/or transcription for in vitro systems, and by cellular expression in in vivo systems.

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In another embodiment, the invention features a method comprising: (a) analyzing the sequence of a RNA target encoded by a target gene; (b) synthesizing one or more sets of siNA molecules having sequence complementary to one or more regions of the RNA of (a); and (c) assaying the siNA molecules of (b) under conditions suitable to determine RNAi targets within the target RNA sequence. In one embodiment, the siNA molecules of (b) have strands of a fixed length, for example about 23 nucleotides in length. In another embodiment, the siNA molecules of (b) are of differing length, for example having strands of about 19 to about 25 (e.g., about 19, 20, 21, 22, 23, 24, or 25) nucleotides in length. In one embodiment, the assay can comprise a reconstituted in vitro siNA assay as described herein. In another embodiment, the assay can comprise a cell culture system in which target RNA is expressed. Fragments of target RNA are analyzed for detectable levels of cleavage, for example by gel electrophoresis, northern blot analysis, or RNAse protection assays, to determine the most suitable target site(s) within the target RNA sequence. The target RNA sequence can be obtained as is known in the art, for example, by cloning and/or transcription for in vitro systems, and by expression in in vivo systems.

By "target site" is meant a sequence within a target RNA that is "targeted" for cleavage mediated by a siNA construct which contains sequences within its antisense region that are complementary to the target sequence.

By "detectable level of cleavage" is meant cleavage of target RNA (and formation of cleaved product RNAs) to an extent sufficient to discern cleavage products above the

background of RNAs produced by random degradation of the target RNA. Production of cleavage products from 1-5% of the target RNA is sufficient to detect above the background for most methods of detection.

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In one embodiment, the invention features a composition comprising a siNA molecule of the invention, which can be chemically-modified, in a pharmaceutically acceptable carrier or diluent. In another embodiment, the invention features a pharmaceutical composition comprising siNA molecules of the invention, which can be chemically-modified, targeting one or more genes in a pharmaceutically acceptable carrier or diluent. In another embodiment, the invention features a method for treating or preventing a disease or condition in a subject, comprising administering to the subject a composition of the invention under conditions suitable for the treatment or prevention of the disease or condition in the subject, alone or in conjunction with one or more other therapeutic compounds. In yet another embodiment, the invention features a method for reducing or preventing tissue rejection in a subject comprising administering to the subject a composition of the invention under conditions suitable for the reduction or prevention of tissue rejection in the subject.

In another embodiment, the invention features a method for validating a gene target, comprising: (a) synthesizing a siNA molecule of the invention, which can be chemically-modified, wherein one of the siNA strands includes a sequence complementary to RNA of a target gene; (b) introducing the siNA molecule into a cell, tissue, or organism under conditions suitable for modulating expression of the target gene in the cell, tissue, or organism; and (c) determining the function of the gene by assaying for any phenotypic change in the cell, tissue, or organism.

In another embodiment, the invention features a method for validating a target gene comprising: (a) synthesizing a siNA molecule of the invention, which can be chemically-modified, wherein one of the siNA strands includes a sequence complementary to RNA of a target gene; (b) introducing the siNA molecule into a biological system under conditions suitable for modulating expression of the target gene in the biological system; and (c) determining the function of the gene by assaying for any phenotypic change in the biological system.

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By "biological system" is meant, material, in a purified or unpurified form, from biological sources, including but not limited to human, animal, plant, insect, bacterial, viral or other sources, wherein the system comprises the components required for RNAi acitivity. The term "biological system" includes, for example, a cell, tissue, or organism, or extract thereof. The term biological system also includes reconstituted RNAi systems that can be used in an *in vitro* setting.

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By "phenotypic change" is meant any detectable change to a cell that occurs in response to contact or treatment with a nucleic acid molecule of the invention (e.g., siNA). Such detectable changes include, but are not limited to, changes in shape, size, proliferation, motility, protein expression or RNA expression or other physical or chemical changes as can be assayed by methods known in the art. The detectable change can also include expression of reporter genes/molecules such as Green Florescent Protein (GFP) or various tags that are used to identify an expressed protein or any other cellular component that can be assayed.

In one embodiment, the invention features a kit containing a siNA molecule of the invention, which can be chemically-modified, that can be used to modulate the expression of a target gene in a cell, tissue, or organism. In another embodiment, the invention features a kit containing more than one siNA molecule of the invention, which can be chemically-modified, that can be used to modulate the expression of more than one target gene in a cell, tissue, or organism.

In one embodiment, the invention features a kit containing a siNA molecule of the invention, which can be chemically-modified, that can be used to modulate the expression of a target gene in a biological system. In another embodiment, the invention features a kit containing more than one siNA molecule of the invention, which can be chemically-modified, that can be used to modulate the expression of more than one target gene in a biological system.

In one embodiment, the invention features a cell containing one or more siNA molecules of the invention, which can be chemically-modified. In another embodiment, the cell containing a siNA molecule of the invention is a mammalian cell. In yet another embodiment, the cell containing a siNA molecule of the invention is a human cell.

In one embodiment, the synthesis of a siNA molecule of the invention, which can be chemically-modified, comprises: (a) synthesis of two complementary strands of the siNA molecule; (b) annealing the two complementary strands together under conditions suitable to obtain a double-stranded siNA molecule. In another embodiment, synthesis of the two complementary strands of the siNA molecule is by solid phase oligonucleotide synthesis. In yet another embodiment, synthesis of the two complementary strands of the siNA molecule is by solid phase tandem oligonucleotide synthesis.

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In one embodiment, the invention features a method for synthesizing a siNA duplex molecule comprising: (a) synthesizing a first oligonucleotide sequence strand of the siNA molecule, wherein the first oligonucleotide sequence strand comprises a cleavable linker molecule that can be used as a scaffold for the synthesis of the second oligonucleotide sequence strand of the siNA; (b) synthesizing the second oligonucleotide sequence strand of siNA on the scaffold of the first oligonucleotide sequence strand, wherein the second oligonucleotide sequence strand further comprises a chemical moiety than can be used to purify the siNA duplex; (c) cleaving the linker molecule of (a) under conditions suitable for the two siNA oligonucleotide strands to hybridize and form a stable duplex; and (d) purifying the siNA duplex utilizing the chemical moiety of the second oligonucleotide sequence strand. In one embodiment, cleavage of the linker molecule in (c) above takes place during deprotection of the oligonucleotide, for example under hydrolysis conditions using an alkylamine base such as methylamine. In one embodiment, the method of synthesis comprises solid phase synthesis on a solid support such as controlled pore glass (CPG) or polystyrene, wherein the first sequence of (a) is synthesized on a cleavable linker, such as a succinyl linker, using the solid support as a scaffold. The cleavable linker in (a) used as a scaffold for synthesizing the second strand can comprise similar reactivity as the solid support derivatized linker, such that cleavage of the solid support derivatized linker and the cleavable linker of (a) takes place concomitantly. In another embodiment, the chemical moiety of (b) that can be used to isolate the attached oligonucleotide sequence comprises a trityl group, for example a dimethoxytrityl group, which can be employed in a trityl-on synthesis strategy as described herein. In yet another embodiment, the chemical moiety, such as a dimethoxytrityl group, is removed during purification, for example, using acidic conditions.

In a further embodiment, the method for siNA synthesis is a solution phase synthesis or hybrid phase synthesis wherein both strands of the siNA duplex are synthesized in tandem using a cleavable linker attached to the first sequence which acts a scaffold for synthesis of the second sequence. Cleavage of the linker under conditions suitable for hybridization of the separate siNA sequence strands results in formation of the double-stranded siNA molecule.

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In another embodiment, the invention features a method for synthesizing a siNA duplex molecule comprising: (a) synthesizing one oligonucleotide sequence strand of the siNA molecule, wherein the sequence comprises a cleavable linker molecule that can be used as a scaffold for the synthesis of another oligonucleotide sequence; (b) synthesizing a second oligonucleotide sequence having complementarity to the first sequence strand on the scaffold of (a), wherein the second sequence comprises the other strand of the doublestranded siNA molecule and wherein the second sequence further comprises a chemical moiety than can be used to isolate the attached oligonucleotide sequence; (c) purifying the product of (b) utilizing the chemical moiety of the second oligonucleotide sequence strand under conditions suitable for isolating the full-length sequence comprising both siNA oligonucleotide strands connected by the cleavable linker and under conditions suitable for the two siNA oligonucleotide strands to hybridize and form a stable duplex. In one embodiment, cleavage of the linker molecule in (c) above takes place during deprotection of the oligonucleotide, for example under hydrolysis conditions. In another embodiment, cleavage of the linker molecule in (c) above takes place after deprotection of the oligonucleotide. In another embodiment, the method of synthesis comprises solid phase synthesis on a solid support such as controlled pore glass (CPG) or polystyrene, wherein the first sequence of (a) is synthesized on a cleavable linker, such as a succinyl linker, using the solid support as a scaffold. The cleavable linker in (a) used as a scaffold for synthesizing the second strand can comprise similar reactivity or differing reactivity as the solid support derivatized linker, such that cleavage of the solid support derivatized linker and the cleavable linker of (a) takes place either concomitantly or sequentially. In one embodiment, the chemical moiety of (b) that can be used to isolate the attached oligonucleotide sequence comprises a trityl group, for example a dimethoxytrityl group.

In another embodiment, the invention features a method for making a doublestranded siNA molecule in a single synthetic process comprising: (a) synthesizing an

oligonucleotide having a first and a second sequence, wherein the first sequence is complementary to the second sequence, and the first oligonucleotide sequence is linked to the second sequence via a cleavable linker, and wherein a terminal 5'-protecting group, for example, a 5'-O-dimethoxytrityl group (5'-O-DMT) remains on the oligonucleotide having the second sequence; (b) deprotecting the oligonucleotide whereby the deprotection results in the cleavage of the linker joining the two oligonucleotide sequences; and (c) purifying the product of (b) under conditions suitable for isolating the double-stranded siNA molecule, for example using a trityl-on synthesis strategy as described herein.

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In another embodiment, the method of synthesis of siNA molecules of the invention comprises the teachings of Scaringe *et al.*, US Patent Nos. 5,889,136; 6,008,400; and 6,111,086, incorporated by reference herein in their entirety.

In one embodiment, the invention features siNA constructs that mediate RNAi in a cell or reconstituted system, wherein the siNA construct comprises one or more chemical modifications, for example, one or more chemical modifications having any of Formulae I-VII or any combination thereof that increases the nuclease resistance of the siNA construct.

In another embodiment, the invention features a method for generating siNA molecules with increased nuclease resistance comprising (a) introducing nucleotides having any of Formula I-VII or any combination thereof into a siNA molecule, and (b) assaying the siNA molecule of step (a) under conditions suitable for isolating siNA molecules having increased nuclease resistance.

In one embodiment, the invention features siNA constructs that mediate RNAi against a target gene, wherein the siNA construct comprises one or more chemical modifications described herein that modulates the binding affinity between the sense and antisense strands of the siNA construct.

In another embodiment, the invention features a method for generating siNA molecules with increased binding affinity between the sense and antisense strands of the siNA molecule comprising (a) introducing nucleotides having any of Formula I-VII or any combination thereof into a siNA molecule, and (b) assaying the siNA molecule of

step (a) under conditions suitable for isolating siNA molecules having increased binding affinity between the sense and antisense strands of the siNA molecule.

In one embodiment, the invention features siNA constructs that mediate RNAi in a cell or reconstituted system, wherein the siNA construct comprises one or more chemical modifications described herein that modulates the binding affinity between the antisense strand of the siNA construct and a complementary target RNA sequence within a cell.

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In one embodiment, the invention features siNA constructs that mediate RNAi in a cell or reconstituted system, wherein the siNA construct comprises one or more chemical modifications described herein that modulates the binding affinity between the antisense strand of the siNA construct and a complementary target DNA sequence within a cell.

In another embodiment, the invention features a method for generating siNA molecules with increased binding affinity between the antisense strand of the siNA molecule and a complementary target RNA sequence comprising (a) introducing nucleotides having any of Formula I-VII or any combination thereof into a siNA molecule, and (b) assaying the siNA molecule of step (a) under conditions suitable for isolating siNA molecules having increased binding affinity between the antisense strand of the siNA molecule and a complementary target RNA sequence.

In another embodiment, the invention features a method for generating siNA molecules with increased binding affinity between the antisense strand of the siNA molecule and a complementary target DNA sequence comprising (a) introducing nucleotides having any of Formula I-VII or any combination thereof into a siNA molecule, and (b) assaying the siNA molecule of step (a) under conditions suitable for isolating siNA molecules having increased binding affinity between the antisense strand of the siNA molecule and a complementary target DNA sequence.

In one embodiment, the invention features siNA constructs that mediate RNAi in a cell or reconstituted system, wherein the siNA construct comprises one or more chemical modifications described herein that modulate the polymerase activity of a cellular polymerase capable of generating additional endogenous siNA molecules having sequence homology to the chemically-modified siNA construct.

In another embodiment, the invention features a method for generating siNA molecules capable of mediating increased polymerase activity of a cellular polymerase capable of generating additional endogenous siNA molecules having sequence homology to a chemically-modified siNA molecule comprising (a) introducing nucleotides having any of Formula I-VII or any combination thereof into a siNA molecule, and (b) assaying the siNA molecule of step (a) under conditions suitable for isolating siNA molecules capable of mediating increased polymerase activity of a cellular polymerase capable of generating additional endogenous siNA molecules having sequence homology to the chemically-modified siNA molecule.

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In one embodiment, the invention features chemically-modified siNA constructs that mediate RNAi in a cell or reconstituted system, wherein the chemical modifications do not significantly effect the interaction of siNA with a target RNA molecule, DNA molecule and/or proteins or other factors that are essential for RNAi in a manner that would decrease the efficacy of RNAi mediated by such siNA constructs.

In another embodiment, the invention features a method for generating siNA molecules with improved RNAi activity, comprising (a) introducing nucleotides having any of Formula I-VII or any combination thereof into a siNA molecule, and (b) assaying the siNA molecule of step (a) under conditions suitable for isolating siNA molecules having improved RNAi activity.

In yet another embodiment, the invention features a method for generating siNA molecules with improved RNAi activity against a target RNA comprising (a) introducing nucleotides having any of Formula I-VII or any combination thereof into a siNA molecule, and (b) assaying the siNA molecule of step (a) under conditions suitable for isolating siNA molecules having improved RNAi activity against the target RNA.

In yet another embodiment, the invention features a method for generating siNA molecules with improved RNAi activity against a DNA target comprising (a) introducing nucleotides having any of Formula I-VII or any combination thereof into a siNA molecule, and (b) assaying the siNA molecule of step (a) under conditions suitable for isolating siNA molecules having improved RNAi activity against the DNA target, such as a gene, chromosome, or portion thereof.

In one embodiment, the invention features siNA constructs that mediate RNAi in a cell or reconstituted system, wherein the siNA construct comprises one or more chemical modifications described herein that modulates the cellular uptake of the siNA construct.

In another embodiment, the invention features a method for generating siNA molecules against a target gene with improved cellular uptake comprising (a) introducing nucleotides having any of Formula I-VII or any combination thereof into a siNA molecule, and (b) assaying the siNA molecule of step (a) under conditions suitable for isolating siNA molecules having improved cellular uptake.

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In one embodiment, the invention features siNA constructs that mediate RNAi against a target gene, wherein the siNA construct comprises one or more chemical modifications described herein that increases the bioavailability of the siNA construct, for example, by attaching polymeric conjugates such as polyethyleneglycol or equivalent conjugates that improve the pharmacokinetics of the siNA construct, or by attaching conjugates that target specific tissue types or cell types *in vivo*. Non-limiting examples of such conjugates are described in Vargeese *et al.*, U.S. Serial No. 10/201,394 incorporated by reference herein.

In one embodiment, the invention features a method for generating siNA molecules of the invention with improved bioavailability, comprising (a) introducing a conjugate into the structure of a siNA molecule, and (b) assaying the siNA molecule of step (a) under conditions suitable for isolating siNA molecules having improved bioavailability. Such conjugates can include ligands for cellular receptors, such as peptides derived from naturally occurring protein ligands; protein localization sequences, including cellular ZIP code sequences; antibodies; nucleic acid aptamers; vitamins and other co-factors, such as folate and N-acetylgalactosamine; polymers, such as polyethyleneglycol (PEG); phospholipids; polyamines, such as spermine or spermidine; and others.

In another embodiment, the invention features a method for generating siNA molecules of the invention with improved bioavailability comprising (a) introducing an excipient formulation to a siNA molecule, and (b) assaying the siNA molecule of step (a) under conditions suitable for isolating siNA molecules having improved bioavailability. Such excipients include polymers such as cyclodextrins, lipids, cationic lipids, polyamines, phospholipids, and others.

In another embodiment, the invention features a method for generating siNA molecules of the invention with improved bioavailability comprising (a) introducing nucleotides having any of Formulae I-VII or any combination thereof into a siNA molecule, and (b) assaying the siNA molecule of step (a) under conditions suitable for isolating siNA molecules having improved bioavailability.

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In another embodiment, polyethylene glycol (PEG) can be covalently attached to siNA compounds of the present invention. The attached PEG can be any molecular weight, preferably from about 2,000 to about 50,000 daltons (Da).

The present invention can be used alone or as a component of a kit having at least one of the reagents necessary to carry out the *in vitro* or *in vivo* introduction of RNA to test samples and/or subjects. For example, preferred components of the kit include a siNA molecule of the invention and a vehicle that promotes introduction of the siNA into cells of interest as described herein (e.g., using lipids and other methods of transfection known in the art, see for example Beigelman *et al*, US 6,395,713). The kit can be used for target validation, such as in determining gene function and/or activity, or in drug optimization, and in drug discovery (see for example Usman et al., USSN 60/402,996). Such a kit can also include instructions to allow a user of the kit to practice the invention.

The term "short interfering nucleic acid", "siNA", "short interfering RNA", "siRNA", "short interfering nucleic acid molecule", "short interfering oligonucleotide molecule", or "chemically-modified short interfering nucleic acid molecule" as used herein refers to any nucleic acid molecule capable of inhibiting or down regulating gene expression or viral replication, for example by mediating RNA interference "RNAi" or gene silencing in a sequence-specific manner; see for example Bass, 2001, Nature, 411, 428-429; Elbashir et al., 2001, Nature, 411, 494-498; and Kreutzer et al., International PCT Publication No. WO 00/44895; Zernicka-Goetz et al., International PCT Publication No. WO 99/32619; Plaetinck et al., International PCT Publication No. WO 00/01846; Mello and Fire, International PCT Publication No. WO 00/01846; Mello and Fire, International PCT Publication No. WO 99/07409; and Li et al., International PCT Publication No. WO 00/44914; Allshire, 2002, Science, 297, 1818-1819; Volpe et al., 2002, Science, 297, 1833-1837; Jenuwein, 2002, Science, 297, 2215-2218; and Hall et al., 2002, Science, 297, 2232-2237;

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Hutvagner and Zamore, 2002, Science, 297, 2056-60; McManus et al., 2002, RNA, 8, 842-850; Reinhart et al., 2002, Gene & Dev., 16, 1616-1626; and Reinhart & Bartel, 2002, Science, 297, 1831). Non limiting examples of siNA molecules of the invention are shown in Figures 4-6, and Tables II, III, and IV herein. For example the siNA can be a double-stranded polynucleotide molecule comprising self-complementary sense and antisense regions, wherein the antisense region comprises nucleotide sequence that is complementary to nucleotide sequence in a target nucleic acid molecule or a portion thereof and the sense region having nucleotide sequence corresponding to the target nucleic acid sequence or a portion thereof. The siNA can be assembled from two separate oligonucleotides, where one strand is the sense strand and the other is the antisense strand, wherein the antisense and sense strands are self-complementary (i.e. each strand comprises nucleotide sequence that is complementary to nucleotide sequence in the other strand; such as where the antisense strand and sense strand form a duplex or double stranded structure, for example wherein the double stranded region is about 19 base pairs): the antisense strand comprises nucleotide sequence that is complementary to nucleotide sequence in a target nucleic acid molecule or a portion thereof and the sense strand comprises nucleotide sequence corresponding to the target nucleic acid sequence or a portion thereof. Alternatively, the siNA is assembled from a single oligonucleotide, where the self-complementary sense and antisense regions of the siNA are linked by means of a nucleic acid based or non-nucleic acid-based linker(s). The siNA can be a polynucleotide with a hairpin secondary structure, having self-complementary sense and antisense regions, wherein the antisense region comprises nucleotide sequence that is complementary to nucleotide sequence in a separate target nucleic acid molecule or a portion thereof and the sense region having nucleotide sequence corresponding to the target nucleic acid sequence or a portion thereof. The siNA can be a circular singlestranded polynucleotide having two or more loop structures and a stem comprising selfcomplementary sense and antisense regions, wherein the antisense region comprises nucleotide sequence that is complementary to nucleotide sequence in a target nucleic acid molecule or a portion thereof and the sense region having nucleotide sequence corresponding to the target nucleic acid sequence or a portion thereof, and wherein the circular polynucleotide can be processed either in vivo or in vitro to generate an active siNA molecule capable of mediating RNAi. The siNA can also comprise a single stranded polynucleotide having nucleotide sequence complementary to nucleotide

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sequence in a target nucleic acid molecule or a portion thereof (for example, where such siNA molecule does not require the presence within the siNA molecule of nucleotide sequence corresponding to the target nucleic acid sequence or a portion thereof), wherein the single stranded polynucleotide can further comprise a terminal phosphate group, such as a 5'-phosphate (see for example Martinez et al., 2002, Cell., 110, 563-574 and Schwarz et al., 2002, Molecular Cell, 10, 537-568), or 5',3'-diphosphate. In certain embodiment, the siNA molecule of the invention comprises separate sense and antisense sequences or regions, wherein the sense and antisense regions are covalently linked by nucleotide or non-nucleotide linkers molecules as is known in the art, or are alternately non-covalently linked by ionic interactions, hydrogen bonding, van der waals interactions, hydrophobic intercations, and/or stacking interactions. In certain embodiments, the siNA molecules of the invention comprise nucleotide sequence that is complementary to nucleotide sequence of a target gene. In another embodiment, the siNA molecule of the invention interacts with nucleotide sequence of a target gene in a manner that causes inhibition of expression of the target gene. As used herein, siNA molecules need not be limited to those molecules containing only RNA, but further encompasses chemically-modified nucleotides and non-nucleotides. In certain embodiments, the short interfering nucleic acid molecules of the invention lack 2'hydroxy (2'-OH) containing nucleotides. Applicant describes in certain embodiments short interfering nucleic acids that do not require the presence of nucleotides having a 2'hydroxy group for mediating RNAi and as such, short interfering nucleic acid molecules of the invention optionally do not include any ribonucleotides (e.g., nucleotides having a 2'-OH group). Such siNA molecules that do not require the presence of ribonucleotides within the siNA molecule to support RNAi can however have an attached linker or linkers or other attached or associated groups, moieties, or chains containing one or more nucleotides with 2'-OH groups. Optionally, siNA molecules can comprise ribonucleotides at about 5, 10, 20, 30, 40, or 50% of the nucleotide positions. The modified short interfering nucleic acid molecules of the invention can also be referred to as short interfering modified oligonucleotides "siMON." As used herein, the term siNA is meant to be equivalent to other terms used to describe nucleic acid molecules that are capable of mediating sequence specific RNAi, for example short interfering RNA (siRNA), doublestranded RNA (dsRNA), micro-RNA (miRNA), short hairpin RNA (shRNA), short interfering oligonucleotide, short interfering nucleic acid, short interfering modified

oligonucleotide, chemically-modified siRNA, post-transcriptional gene silencing RNA (ptgsRNA), and others. In addition, as used herein, the term RNAi is meant to be equivalent to other terms used to describe sequence specific RNA interference, such as post transcriptional gene silencing, or epigenetics. For example, siNA molecules of the invention can be used to epigenetically silence genes at both the post-transcriptional level or the pre-transcriptional level. In a non-limiting example, epigenetic regulation of gene expression by siNA molecules of the invention can result from siNA mediated modification of chromatin structure to alter gene expression (see, for example, Allshire, 2002, Science, 297, 1818-1819; Volpe et al., 2002, Science, 297, 1833-1837; Jenuwein, 2002, Science, 297, 2215-2218; and Hall et al., 2002, Science, 297, 2232-2237).

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By "modulate" is meant that the expression of the gene, or level of RNA molecule or equivalent RNA molecules encoding one or more proteins or protein subunits, or activity of one or more proteins or protein subunits is up regulated or down regulated, such that expression, level, or activity is greater than or less than that observed in the absence of the modulator. For example, the term "modulate" can mean "inhibit," but the use of the word "modulate" is not limited to this definition.

By "inhibit" it is meant that the activity of a gene expression product or level of RNAs or equivalent RNAs encoding one or more gene products is reduced below that observed in the absence of the nucleic acid molecule of the invention. In one embodiment, inhibition with a siNA molecule preferably is below that level observed in the presence of an inactive or attenuated molecule that is unable to mediate an RNAi response. In another embodiment, inhibition of gene expression with the siNA molecule of the instant invention is greater in the presence of the siNA molecule than in its absence.

By "inhibit", "down-regulate", or "reduce", it is meant that the expression of the gene, or level of RNA molecules or equivalent RNA molecules encoding one or more proteins or protein subunits, or activity of one or more proteins or protein subunits, is reduced below that observed in the absence of the nucleic acid molecules (e.g., siNA) of the invention. In one embodiment, inhibition, down-regulation or reduction with an siNA molecule is below that level observed in the presence of an inactive or attenuated molecule. In another embodiment, inhibition, down-regulation, or reduction with siNA

molecules is below that level observed in the presence of, for example, an siNA molecule with scrambled sequence or with mismatches. In another embodiment, inhibition, down-regulation, or reduction of gene expression with a nucleic acid molecule of the instant invention is greater in the presence of the nucleic acid molecule than in its absence.

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By "gene" or "target gene" is meant, a nucleic acid that encodes an RNA, for example, nucleic acid sequences including, but not limited to, structural genes encoding a polypeptide. The target gene can be a gene derived from a cell, an endogenous gene, a transgene, or exogenous genes such as genes of a pathogen, for example a virus, which is present in the cell after infection thereof. The cell containing the target gene can be derived from or contained in any organism, for example a plant, animal, protozoan, virus, bacterium, or fungus. Non-limiting examples of plants include monocots, dicots, or gymnosperms. Non-limiting examples of animals include vertebrates or invertebrates. Non-limiting examples of fungi include molds or yeasts.

By "endogenous" or "cellular" gene is meant a gene normally found in a cell in its natural location in the genome. For example, HER-2, VEGF, VEGF-R, EGFR, BCL-2, c-MYC, RAS and the like would be considered an endogenous gene. Genes expressed in a cell from a plasmid, viral vector or other vectors or from virus, bacteria, fungi would be considered "foreign" or "heterologous" gene; such genes are not normally found in the host cell, but are introduced by standard gene transfer techniques or as a result of infection by a virus, bacterial or other infectious agent.

By "gene family" is meant a group of more than one nucleic acid molecules that share at least one common characteristic, such as sequence homology, target specificity, mode of action, secondary structure, or the ability to modulate a process or more than one process in a biological system. The gene family can be of viral or cellular origin. The gene family can encode, for example, groups of cytokines, receptors, growth factors, adapter proteins, structural proteins, and other protein epitopes.

By "protein family" is meant a group of more than one proteins, peptides, or polypeptides that share at least one common characteristic, such as sequence homology, target specificity, mode of action, secondary structure, or the ability to modulate a process or more than one process in a biological system. The protein family can be of viral or

cellular origin. The protein family can encode, for example, groups of cytokines, receptors, growth factors, adapter proteins, structural proteins, and other protein epitopes.

By "highly conserved sequence region" is meant, a nucleotide sequence of one or more regions in a target gene does not vary significantly from one generation to the other or from one biological system to the other.

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By "cancer" is meant a group of diseases characterized by uncontrolled growth and spread of abnormal cells.

By "sense region" is meant a nucleotide sequence of a siNA molecule having complementarity to an antisense region of the siNA molecule. In addition, the sense region of a siNA molecule can comprise a nucleic acid sequence having homology with a target nucleic acid sequence.

By "antisense region" is meant a nucleotide sequence of a siNA molecule having complementarity to a target nucleic acid sequence. In addition, the antisense region of a siNA molecule can optionally comprise a nucleic acid sequence having complementarity to a sense region of the siNA molecule.

By "target nucleic acid" is meant any nucleic acid sequence whose expression or activity is to be modulated. The target nucleic acid can be DNA or RNA.

By "complementarity" is meant that a nucleic acid can form hydrogen bond(s) with another nucleic acid sequence by either traditional Watson-Crick or other non-traditional types. In reference to the nucleic molecules of the present invention, the binding free energy for a nucleic acid molecule with its complementary sequence is sufficient to allow the relevant function of the nucleic acid to proceed, e.g., RNAi activity. Determination of binding free energies for nucleic acid molecules is well known in the art (see, e.g., Turner et al., 1987, CSH Symp. Quant. Biol. LII pp.123-133; Frier et al., 1986, Proc. Nat. Acad. Sci. USA 83:9373-9377; Turner et al., 1987, J. Am. Chem. Soc. 109:3783-3785). A percent complementarity indicates the percentage of contiguous residues in a nucleic acid molecule that can form hydrogen bonds (e.g., Watson-Crick base pairing) with a second nucleic acid sequence (e.g., 5, 6, 7, 8, 9, 10 out of 10 being 50%, 60%, 70%, 80%, 90%, and 100% complementary). "Perfectly complementary" means that all the contiguous

residues of a nucleic acid sequence will hydrogen bond with the same number of contiguous residues in a second nucleic acid sequence.

The siNA molecules of the invention represent a novel therapeutic approach to a broad spectrum of diseases and conditions, including cancer or cancerous disease, infectious disease, cardiovascular disease, neurological disease, prion disease, inflammatory disease, autoimmune disease, pulmonary disease, renal disease, liver disease, mitochondrial disease, endocrine disease, reproduction related diseases and conditions, and any other indications that can respond to the level of an expressed gene product in a cell or organsim.

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In one embodiment of the present invention, each sequence of a siNA molecule of the invention is independently about 18 to about 24 nucleotides in length, in specific embodiments about 18, 19, 20, 21, 22, 23, or 24 nucleotides in length. In another embodiment, the siNA duplexes of the invention independently comprise about 17 to about 23 base pairs (e.g., about 17, 18, 19, 20, 21, 22 or 23). In yet another embodiment, siNA molecules of the invention comprising hairpin or circular structures are about 35 to about 55 (e.g., about 35, 40, 45, 50 or 55) nucleotides in length, or about 38 to about 44 (e.g., 38, 39, 40, 41, 42, 43 or 44) nucleotides in length and comprising about 16 to about 22 (e.g., about 16, 17, 18, 19, 20, 21 or 22) base pairs. Exemplary siNA molecules of the invention are shown in **Table II**. Exemplary synthetic siNA molecules of the invention are shown in **Table II** and/or **Figures 18-19**.

As used herein "cell" is used in its usual biological sense, and does not refer to an entire multicellular organism, e.g., specifically does not refer to a human. The cell can be present in an organism, e.g., birds, plants and mammals such as humans, cows, sheep, apes, monkeys, swine, dogs, and cats. The cell can be prokaryotic or eukaryotic (e.g., mammalian or plant cell). The cell can be of somatic or germ line origin, totipotent or pluripotent, dividing or non-dividing. The cell can also be derived from or can comprise a gamete or embryo, a stem cell, or a fully differentiated cell.

The siNA molecules of the invention are added directly, or can be complexed with cationic lipids, packaged within liposomes, or otherwise delivered to target cells or tissues. The nucleic acid or nucleic acid complexes can be locally administered to relevant tissues ex vivo, or in vivo through injection, infusion pump or stent, with or

without their incorporation in biopolymers. In particular embodiments, the nucleic acid molecules of the invention comprise sequences shown in Tables I-II and/or Figures 18-19. Examples of such nucleic acid molecules consist essentially of sequences defined in these tables and figures. Furthermore, the chemically modified constructs described in Table IV can be applied to any siNA sequence of the invention.

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In another aspect, the invention provides mammalian cells containing one or more siNA molecules of this invention. The one or more siNA molecules can independently be targeted to the same or different sites.

By "RNA" is meant a molecule comprising at least one ribonucleotide residue. By "ribonucleotide" is meant a nucleotide with a hydroxyl group at the 2' position of a β-D-ribo-furanose moiety. The terms include double-stranded RNA, single-stranded RNA, isolated RNA such as partially purified RNA, essentially pure RNA, synthetic RNA, recombinantly produced RNA, as well as altered RNA that differs from naturally occurring RNA by the addition, deletion, substitution and/or alteration of one or more nucleotides. Such alterations can include addition of non-nucleotide material, such as to the end(s) of the siNA or internally, for example at one or more nucleotides of the RNA. Nucleotides in the RNA molecules of the instant invention can also comprise non-standard nucleotides, such as non-naturally occurring nucleotides or chemically synthesized nucleotides or deoxynucleotides. These altered RNAs can be referred to as analogs or analogs of naturally-occurring RNA.

By "subject" is meant an organism, which is a donor or recipient of explanted cells or the cells themselves. "Subject" also refers to an organism to which the nucleic acid molecules of the invention can be administered. In one embodiment, a subject is a mammal or mammalian cells. In another embodiment, a subject is a human or human cells.

The term "phosphorothioate" as used herein refers to an internucleotide linkage having Formula I, wherein Z and/or W comprise a sulfur atom. Hence, the term phosphorothioate refers to both phosphorothioate and phosphorodithioate internucleotide linkages.

The term "universal base" as used herein refers to nucleotide base analogs that form base pairs with each of the natural DNA/RNA bases with little discrimination between them. Non-limiting examples of universal bases include C-phenyl, C-naphthyl and other aromatic derivatives, inosine, azole carboxamides, and nitroazole derivatives such as 3-nitropyrrole, 4-nitroindole, 5-nitroindole, and 6-nitroindole as known in the art (see for example Loakes, 2001, *Nucleic Acids Research*, 29, 2437-2447).

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The term "acyclic nucleotide" as used herein refers to any nucleotide having an acyclic ribose sugar, for example where any of the ribose carbons (C1, C2, C3, C4, or C5), are independently or in combination absent from the nucleotide.

The nucleic acid molecules of the instant invention, individually, or in combination or in conjunction with other drugs, can be used to treat diseases or conditions discussed herein. For example, to treat a particular disease or condition, the siNA molecules can be administered to a subject or can be administered to other appropriate cells evident to those skilled in the art, individually or in combination with one or more drugs under conditions suitable for the treatment.

In a further embodiment, the siNA molecules can be used in combination with other known treatments to treat conditions or diseases discussed above. For example, the described molecules could be used in combination with one or more known therapeutic agents to treat a disease or condition. Non-limiting examples of other therapeutic agents that can be readily combined with a siNA molecule of the invention are enzymatic nucleic acid molecules, allosteric nucleic acid molecules, antisense, decoy, or aptamer nucleic acid molecules, antibodies such as monoclonal antibodies, small molecules, and other organic and/or inorganic compounds including metals, salts and ions.

In one embodiment, the invention features an expression vector comprising a nucleic acid sequence encoding at least one siNA molecule of the invention, in a manner which allows expression of the siNA molecule. For example, the vector can contain sequence(s) encoding both strands of a siNA molecule comprising a duplex. The vector can also contain sequence(s) encoding a single nucleic acid molecule that is self-complementary and thus forms a siNA molecule. Non-limiting examples of such expression vectors are described in Paul et al., 2002, Nature Biotechnology, 19, 505; Miyagishi and Taira, 2002, Nature Biotechnology, 19, 497; Lee et al., 2002, Nature

Biotechnology, 19, 500; and Novina et al., 2002, Nature Medicine, advance online publication doi:10.1038/nm725.

In another embodiment, the invention features a mammalian cell, for example, a human cell, including an expression vector of the invention.

In yet another embodiment, the expression vector of the invention comprises a sequence for a siRNA molecule having complementarity to a RNA molecule referred to by a Genbank Accession number in Table III.

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In yet another embodiment, the expression vector of the invention comprises a sequence for a siNA molecule having complementarity to a RNA molecule referred to by a Genbank Accession numbers, for example Genbank Accession Nos. shown in **Table I**.

In one embodiment, an expression vector of the invention comprises a nucleic acid sequence encoding two or more siNA molecules, which can be the same or different.

In another aspect of the invention, siRNA molecules that interact with target RNA molecules and down-regulate gene encoding target RNA molecules (for example target RNA molecules referred to by Genbank Accession number in Table III) are expressed from transcription units inserted into DNA or RNA vectors. The recombinant vectors can be DNA plasmids or viral vectors siNA expressing viral vectors can be constructed based on, but not limited to, adeno-associated virus, retrovirus, adenovirus, or alphavirus. The recombinant vectors capable of expressing the siNA molecules can be delivered as described herein, and persist in target cells. Alternatively, viral vectors can be used that provide for transient expression of siNA molecules. Such vectors can be repeatedly administered as necessary. Once expressed, the siNA molecules bind and down-regulate gene function or expression via RNA interference (RNAi). Delivery of siNA expressing vectors can be systemic, such as by intravenous or intramuscular administration, by administration to target cells ex-planted from a subject followed by reintroduction into the subject, or by any other means that would allow for introduction into the desired target cell.

By "vectors" is meant any nucleic acid- and/or viral-based technique used to deliver a desired nucleic acid.

Other features and advantages of the invention will be apparent from the following description of the preferred embodiments thereof, and from the claims.

BRIEF DESCRIPTION OF THE DRAWINGS

Figure 1 shows a non-limiting example of a scheme for the synthesis of siNA molecules. The complementary siNA sequence strands, strand 1 and strand 2, are synthesized in tandem and are connected by a cleavable linkage, such as a nucleotide succinate or abasic succinate, which can be the same or different from the cleavable linker used for solid phase synthesis on a solid support. The synthesis can be either solid phase or solution phase, in the example shown, the synthesis is a solid phase synthesis. The synthesis is performed such that a protecting group, such as a dimethoxytrityl group, remains intact on the terminal nucleotide of the tandem oligonucleotide. Upon cleavage and deprotection of the oligonucleotide, the two siNA strands spontaneously hybridize to form a siNA duplex, which allows the purification of the duplex by utilizing the properties of the terminal protecting group, for example by applying a trityl on purification method wherein only duplexes/oligonucleotides with the terminal protecting group are isolated.

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Figure 2 shows a MALDI-TOV mass spectrum of a purified siNA duplex synthesized by a method of the invention. The two peaks shown correspond to the predicted mass of the separate siNA sequence strands. This result demonstrates that the siNA duplex generated from tandem synthesis can be purified as a single entity using a simple trityl-on purification methodology.

Figure 3 shows the results of a stability assay used to determine the serum stability of chemically modified siNA constructs compared to a siNA control consisting of all RNA with 3'-TT termini. T ½ values are shown for duplex stability.

Figure 4 shows the results of an RNAi activity screen of phosphorothioate modified siNA constructs using a luciferase reporter system.

Figure 5 shows the results of an RNAi activity screen of phosphorothioate and universal base modified siNA constructs using a luciferase reporter system.

Figure 6 shows the results of an RNAi activity screen of 2'-O-methyl modified siNA constructs using a luciferase reporter system.

Figure 7 shows the results of an RNAi activity screen of 2'-O-methyl and 2'-deoxy-2'-fluoro modified siNA constructs using a luciferase reporter system.

Figure 8 shows the results of an RNAi activity screen of a phosphorothioate modified siNA construct using a luciferase reporter system.

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Figure 9 shows the results of an RNAi activity screen of an inverted deoxyabasic modified siNA construct generated via tandem synthesis using a luciferase reporter system.

Figure 10 shows the results of an RNAi activity screen of chemically modified siNA constructs including 3'-glyceryl modified siNA constructs compared to an all RNA control siNA construct using a luciferase reporter system. These chemically modified siNAs were compared in the luciferase assay described herein at 1 nM and 10nM concentration using an all RNA siNA control (siGL2) having having 3'-terminal dithymidine (TT) and its corresponding inverted control (Inv siGL2). The background level of luciferase expression in the HeLa cells is designated by the "cells" column. Sense and antisense strands of chemically modified siNA constructs are shown by RPI number (sense strand/antisense strand). Sequences correspoding to these RPI numbers are shown in Table I.

Figure 11 shows the results of an RNAi activity screen of chemically modified siNA constructs. The screen compared various combinations of sense strand chemical modifications and antisense strand chemical modifications. These chemically modified siNAs were compared in the luciferase assay described herein at 1 nM and 10nM concentration using an all RNA siNA control (siGL2) having having 3'-terminal dithymidine (TT) and its corresponding inverted control (Inv siGL2). The background level of luciferase expression in the HeLa cells is designated by the "cells" column. Sense and antisense strands of chemically modified siNA constructs are shown by RPI number (sense strand/antisense strand). Sequences correspoding to these RPI numbers are shown in Table I.

Figure 12 shows the results of an RNAi activity screen of chemically modified siNA constructs. The screen compared various combinations of sense strand chemical modifications and antisense strand chemical modifications. These chemically modified siNAs were compared in the luciferase assay described herein at 1 nM and 10nM concentration using an all RNA siNA control (siGL2) having having 3'-terminal dithymidine (TT) and its corresponding inverted control (Inv siGL2). The background level of luciferase expression in the HeLa cells is designated by the "cells" column. Sense and antisense strands of chemically modified siNA constructs are shown by RPI number (sense strand/antisense strand). Sequences corresponding to these RPI numbers are shown in Table I. In addition, the antisense strand alone (RPI 30430) and an inverted control (RPI 30227/30229, having matched chemistry to RPI 30063/30224) was compared to the siNA duplexes described above.

Figure 13 shows the results of an RNAi activity screen of chemically modified siNA constructs. The screen compared various combinations of sense strand chemical modifications and antisense strand chemical modifications. These chemically modified siNAs were compared in the luciferase assay described herein at 1 nM and 10nM concentration using an all RNA siNA control (siGL2) having having 3'-terminal dithymidine (TT) and its corresponding inverted control (Inv siGL2). The background level of luciferase expression in the HeLa cells is designated by the "cells" column. Sense and antisense strands of chemically modified siNA constructs are shown by RPI number (sense strand/antisense strand). Sequences correspoding to these RPI numbers are shown in Table I. In addition, an inverted control (RPI 30226/30229, having matched chemistry to RPI 30222/30224) was compared to the siNA duplexes described above.

Figure 14 shows the results of an RNAi activity screen of chemically modified siNA constructs including various 3'-terminal modified siNA constructs compared to an all RNA control siNA construct using a luciferase reporter system. These chemically modified siNAs were compared in the luciferase assay described herein at 1 nM and 10nM concentration using an all RNA siNA control (siGL2) having having 3'-terminal dithymidine (TT) and its corresponding inverted control (Inv siGL2). The background level of luciferase expression in the HeLa cells is designated by the "cells" column. Sense and antisense strands of chemically modified siNA constructs are shown by RPI

number (sense strand/antisense strand). Sequences correspoding to these RPI numbers are shown in Table I.

Figure 15 shows the results of an RNAi activity screen of chemically modified siNA constructs. The screen compared various combinations of sense strand chemistries compared to a fixed antisense strand chemistry. These chemically modified siNAs were compared in the luciferase assay described herein at 1 nM and 10nM concentration using an all RNA siNA control (siGL2) having having 3'-terminal dithymidine (TT) and its corresponding inverted control (Inv siGL2). The background level of luciferase expression in the HeLa cells is designated by the "cells" column. Sense and antisense strands of chemically modified siNA constructs are shown by RPI number (sense strand/antisense strand). Sequences correspoding to these RPI numbers are shown in Table I.

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Figure 16 shows the results of a siNA titration study wherein the RNAi activity of a phosphorothicate modified siNA construct is compared to that of a siNA construct consisting of all ribonucleotides except for two terminal thymidine residues using a luciferase reporter system.

Figure 17 shows a non-limiting proposed mechanistic representation of target RNA degradation involved in RNAi. Double-stranded RNA (dsRNA), which is generated by RNA-dependent RNA polymerase (RdRP) from foreign single-stranded RNA, for example viral, transposon, or other exogenous RNA, activates the DICER enzyme that in turn generates siNA duplexes. Alternately, synthetic or expressed siNA can be introduced directely into a cell by appropriate means. An active siNA complex forms which recognizes a target RNA, resulting in degradation of the target RNA by the RISC endonuclease complex or in the synthesis of additional RNA by RNA-dependent RNA polymerase (RdRP), which can activate DICER and result in additional siNA molecules, thereby amplifying the RNAi response.

Figure 18A-F shows non-limiting examples of chemically-modified siNA constructs of the present invention. In the figure, N stands for any nucleotide (adenosine, guanosine, cytosine, uridine, or optionally thymidine, for example thymidine can be substituted in the overhanging regions designated by parenthesis (N N). Various modifications are shown for the sense and antisense strands of the siNA constructs.

The sense strand comprises 21 nucleotides having four Figure 18A: phosphorothioate 5'- and 3'-terminal internucleotide linkages, wherein the two terminal 3'nucleotides are optionally base paired and wherein all pyrimidine nucleotides that may be present are 2'-O-methyl or 2'-deoxy-2'-fluoro modified nucleotides except for (N N) nucleotides, which can comprise ribonucleotides, deoxynucleotides, universal bases, or other chemical modifications described herein. The antisense strand comprises 21 nucleotides, optionally having a 3'-terminal glyceryl moiety and wherein the two terminal 3'-nucleotides are optionally complementary to the target RNA sequence, and having one linkage 3'-terminal phosphorothioate internucleotide and four 5'-terminal phosphorothioate internucleotide linkages and wherein all pyrimidine nucleotides that may be present are 2'-deoxy-2'-fluoro modified nucleotides except for (N N) nucleotides, which can comprise ribonucleotides, deoxynucleotides, universal bases, or other chemical modifications described herein.

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Figure 18B: The sense strand comprises 21 nucleotides wherein the two terminal 3'-nucleotides are optionally base paired and wherein all pyrimidine nucleotides that may be present are 2'-O-methyl or 2'-deoxy-2'-fluoro modified nucleotides except for (N N) nucleotides, which can comprise ribonucleotides, deoxynucleotides, universal bases, or other chemical modifications described herein. The antisense strand comprises 21 nucleotides, optionally having a 3'-terminal glyceryl moiety and wherein the two terminal 3'-nucleotides are optionally complementary to the target RNA sequence, and wherein all pyrimidine nucleotides that may be present are 2'-deoxy-2'-fluoro modified nucleotides except for (N N) nucleotides, which can comprise ribonucleotides, deoxynucleotides, universal bases, or other chemical modifications described herein.

Figure 18C: The sense strand comprises 21 nucleotides having 5'- and 3'- terminal cap moieties wherein the two terminal 3'-nucleotides are optionally base paired and wherein all pyrimidine nucleotides that may be present are 2'-O-methyl or 2'-deoxy-2'-fluoro modified nucleotides except for (N N) nucleotides, which can comprise ribonucleotides, deoxynucleotides, universal bases, or other chemical modifications described herein. The antisense strand comprises 21 nucleotides, optionally having a 3'-terminal glyceryl moiety and wherein the two terminal 3'-nucleotides are optionally complementary to the target RNA sequence, and having one 3'-terminal phosphorothioate internucleotide linkage and wherein all pyrimidine nucleotides that may be present are 2'-

deoxy-2'-fluoro modified nucleotides except for (N N) nucleotides, which can comprise ribonucleotides, deoxynucleotides, universal bases, or other chemical modifications described herein.

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Figure 18D: The sense strand comprises 21 nucleotides having 5'- and 3'- terminal cap moieties wherein the two terminal 3'-nucleotides are optionally base paired and wherein all pyrimidine nucleotides that may be present are 2'-deoxy-2'-fluoro modified nucleotides except for (N N) nucleotides, which can comprise ribonucleotides, deoxynucleotides, universal bases, or other chemical modifications described herein and wherein and all purine nucleotides that may be present are 2'-deoxy nucleotides. The antisense strand comprises 21 nucleotides, optionally having a 3'-terminal glyceryl moiety and wherein the two terminal 3'-nucleotides are optionally complementary to the target RNA sequence, and having one 3'-terminal phosphorothioate internucleotide linkage and wherein all pyrimidine nucleotides that may be present are 2'-deoxy-2'-fluoro modified nucleotides and all purine nucleotides that may be present are 2'-O-methyl modified nucleotides except for (N N) nucleotides, which can comprise ribonucleotides, deoxynucleotides, universal bases, or other chemical modifications described herein.

Figure 18E: The sense strand comprises 21 nucleotides having 5'- and 3'- terminal cap moieties wherein the two terminal 3'-nucleotides are optionally base paired and wherein all pyrimidine nucleotides that may be present are 2'-deoxy-2'-fluoro modified nucleotides except for (N N) nucleotides, which can comprise ribonucleotides, deoxynucleotides, universal bases, or other chemical modifications described herein. The antisense strand comprises 21 nucleotides, optionally having a 3'-terminal glyceryl moiety and wherein the two terminal 3'-nucleotides are optionally complementary to the target RNA sequence, and wherein all pyrimidine nucleotides that may be present are 2'-deoxy-2'-fluoro modified nucleotides and all purine nucleotides that may be present are 2'-O-methyl modified nucleotides except for (N N) nucleotides, which can comprise ribonucleotides, deoxynucleotides, universal bases, or other chemical modifications described herein.

Figure 18F: The sense strand comprises 21 nucleotides having 5'- and 3'- terminal cap moieties wherein the two terminal 3'-nucleotides are optionally base paired and wherein all pyrimidine nucleotides that may be present are 2'-deoxy-2'-fluoro modified

nucleotides except for (N N) nucleotides, which can comprise ribonucleotides, deoxynucleotides, universal bases, or other chemical modifications described herein. The antisense strand comprises 21 nucleotides, optionally having a 3'-terminal glyceryl moiety and wherein the two terminal 3'-nucleotides are optionally complementary to the target RNA sequence, and having one 3'-terminal phosphorothioate internucleotide linkage and wherein all pyrimidine nucleotides that may be present are 2'-deoxy-2'-fluoro modified nucleotides and all purine nucleotides that may be present are 2'-deoxy modified nucleotides except for (N N) nucleotides, which can comprise ribonucleotides, deoxynucleotides, universal bases, or other chemical modifications described herein. The antisense strand of constructs A-F comprise sequence complementary to target RNA sequence of the invention.

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Figure 19 shows non-limiting examples of specific chemically modified siNA sequences of the invention. A-F applies the chemical modifications described in Figure 18A-F to a representative siNA sequence targeting the EGFR (HER1).

Figure 20 shows non-limiting examples of different siNA constructs of the invention. The examples shown (constructs 1, 2, and 3) have 19 representative base pairs, however, different embodiments of the invention include any number of base pairs Bracketed regions represent nucleotide overhangs, for example described herein. comprising between about 1, 2, 3, or 4 nucleotides in length, preferably about 2 nucleotides. Constructs 1 and 2 can be used independently for RNAi activity. Construct 2 can comprise a polynucleotide or non-nucleotide linker, which can optionally be designed as a biodegradable linker. In one embodiment, the loop structure shown in construct 2 can comprise a biodegradable linker that results in the formation of construct 1 in vivo and/or in vitro. In another example, construct 3 can be used to generate construct 2 under the same principle wherein a linker is used to generate the active siNA construct 2 in vivo and/or in vitro, which can optionally utilize another biodegradable linker to generate the active siNA construct 1 in vivo and/or in vitro. As such, the stability and/or activity of the siNA constructs can be modulated based on the design of the siNA construct for use in vivo or in vitro and/or in vitro.

Figure 21 is a diagrammatic representation of a method used to determine target sites for siNA mediated RNAi within a particular target nucleic acid sequence, such as

messenger RNA. (A) A pool of siNA oligonucleotides are synthesized wherein the antisense region of the siNA constructs has complementarity to target sites across the target nucleic acid sequence, and wherein the sense region comprises sequence complementary to the antisense region of the siNA. (B) The sequences are transfected into cells. (C) Cells are selected based on phenotypic change that is associated with modulation of the target nucleic acid sequence. (D) The siNA is isolated from the selected cells and is sequenced to identify efficacious target sites within the target nucleic acid sequence.

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Figure 22 shows non-limiting examples of different stabilization chemistries (1-10) that can be used, for example, to stabilize the 3'-end of siNA sequences of the invention, including (1) [3-3']-inverted deoxyribose; (2) deoxyribonucleotide; (3) [5'-3']-3'-deoxyribonucleotide; (4) [5'-3']-ribonucleotide; (5) [5'-3']-3'-O-methyl ribonucleotide; (6) 3'-glyceryl; (7) [3'-5']-3'-deoxyribonucleotide; (8) [3'-3']-deoxyribonucleotide; (9) [5'-2']-deoxyribonucleotide; and (10) [5-3']-dideoxyribonucleotide. In addition to modified and unmodified backbone chemistries indicated in the figure, these chemistries can be combined with different backbone modifications as described herein, for example, backbone modifications having Formula I. In addition, the 2'-deoxy nucleotide shown 5' to the terminal modifications shown can be another modified or unmodified nucleotide or non-nucleotide described herein, for example modifications having any of Formulae I-VII or any combination thereof.

Figure 23 shows a non-limiting example of siNA mediated inhibition of VEGF-induced angiogenesis using the rat corneal model of angiogenesis. siNA targeting site 2340 of VEGFR1 RNA (shown as RPI No. sense strand/antisense strand) were compared to inverted controls (shown as RPI No. sense strand/antisense strand) at three different concentrations and compared to a VEGF control in which no siNA was administered.

Figure 24 shows a non-limiting example of a strategy used to identify chemically modified siNA constructs of the invention that are nuclease resistance while preserving the ability to mediate RNAi activity. Chemical modifications are introduced into the siNA construct based on educated design parameters (e.g. introducing 2'-mofications, base modifications, backbone modifications, terminal cap modifications etc). The modified construct in tested in an appropriate system (e.g human serum for nuclease

resistance, shown, or an animal model for PK/delivery parameters). In parallel, the siNA construct is tested for RNAi activity, for example in a cell culture system such as a luciferase reporter assay). Lead siNA constructs are then identified which possess a particular characteristic while maintaining RNAi activity, and can be further modified and assayed once again. This same approach can be used to identify siNA-conjugate molecules with improved pharmacokinetic profiles, delivery, and RNAi activity.

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Figure 25 shows a non-limiting example of reduction of HER2 mRNA in A549 cells mediated by RNA-based and chemically-modified siNAs that target HER2 mRNA sites 2344 and 3706. A549 cells were transfected with 4 ug/ml lipid complexed with 25 nM unmodified siNA with a 3'-terminal dithymidine cap (RPI#28266/28267) or a corresponding inverted control (RPI#28268/28269) for site 2344 and (RPI#28262/28263) and a corresponding inverted control (RPI 28264/28265) for site 3706. In addition, A549 cells were transfected with 4 ug/ml lipid complexed with 25 nM modified siNA (RPI#30442/30443) and a corresponding matched control (RPI#30444/30445) for site 2344 and (RPI#30438/30439) and a corresponding matched control (RPI 30440/30441) for site 3706. As shown in the figures, the modified and unmodified constructs targeting sites 2344 and 3706 all demonstrate significant inhibition of HER2 RNA expression.

Figure 26 shows a non-limiting example of reduction of PKC-alpha mRNA in A549 cells mediated by chemically-modified siNAs that target PKC-alpha mRNA. A549 cells were transfected with 0.25 ug/well of lipid complexed with 25 nM siNA. A screen of siNA constructs comprising ribonucleotides and 3'-terminal dithymidine caps was compared to untreated cells, scrambled siNA control constructs (Scram1 and Scram2), and cells transfected with lipid alone (transfection control). As shown in the figure, all of the siNA constructs show significant reduction of PKC-alpha RNA expression.

Figure 27 shows a non-limiting example of reduction of Myc (c-Myc) mRNA in 293T cells mediated by chemically-modified siNAs that target c-Myc mRNA. 293T cells were transfected with 0.25 ug/well of lipid complexed with 25 nM siNA. A screen of siNA constructs comprising ribonucleotides and 3'-terminal dithymidine caps was compared to untreated cells, scrambled siNA control constructs (Scram1 and Scram2), and cells transfected with lipid alone (transfection control). As shown in the figure, three

of the siNA constructs (RPI 30993/31069; RPI 30995/31071; and RPI 30996/31072) show significant reduction of c-Myc RNA expression.

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Figure 28 shows a non-limiting example of reduction of BCL2 mRNA in A549 cells mediated by chemically-modified siNAs that target BCL2 mRNA. A549 cells were transfected with 0.25 ug/well of lipid complexed with 25 nM siNA. A siNA construct comprising ribonucleotides and 3'-terminal dithymidine caps (RPI#30998/31074) was tested along with a chemically modified siNA construct comprising 2'-deoxy-2'-fluoro pyrimidine nucleotides and purine ribonucleotides in which the sense strand of the siNA is further modified with 5' and 3'-terminal inverted deoxyabasic caps and the antisense strand comprises a 3'-terminal phosphorothioate internucleotide linkage (RPI#31368/31369), which was also compared to a matched chemistry inverted control (RPI#31370/31371) and a chemically modified siNA construct comprising 2'-deoxy-2'fluoro pyrimidine and 2'-deoxy-2'-fluoro purine nucleotides in which the sense strand of the siNA is further modified with 5' and 3'-terminal inverted deoxyabasic caps and the antisense strand comprises a 3'-terminal phosphorothioate internucleotide linkage (RPI#31372/31373) which was also compared to a matched chemistry inverted control (RPI#31374/31375). In addition, the siNA constructs were also compared to untreated cells, cells transfected with lipid and scrambled siNA constructs (Scram1 and Scram2), and cells transfected with lipid alone (transfection control). As shown in the figure, the siNA constructs show significant reduction of BCL2 RNA expression compared to scrambled, untreated, and transfection controls.

Figure 29 shows a non-limiting example of reduction of CHK-1 mRNA in A549 cells mediated by chemically-modified siNAs that target CHK-1 mRNA. A549 cells were transfected with 0.25 ug/well of lipid complexed with 25 nM siNA. A siNA construct comprising ribonucleotides and 3'-terminal dithymidine caps (RPI#31003/31079) and a chemically modified siNA construct comprising 2'-deoxy-2'fluoro pyrimidine nucleotides and purine ribonucleotides in which the sense strand of the siNA is further modified with 5' and 3'-terminal inverted deoxyabasic caps and in which the antisense strand comprises a 3'-terminal phosphorothioate internucleotide linkage (RPI#31302/31303), were compared to a matched chemistry inverted control (RPI#31314/31325). In addition, the siNA constructs were also compared to untreated cells, cells transfected with lipid and scrambled siNA constructs (Scram1 and Scram2),

and cells transfected with lipid alone (transfection control). As shown in the figure, both siNA constructs show significant reduction of CHK-1 RNA expression compared to appropriate controls.

Figure 30 shows a non-limiting example of reduction of BACE mRNA in A549 cells mediated by siNAs that target BACE mRNA. A549 cells were transfected with 0.25 ug/well of lipid complexed with 25 nM siNA. A screen of siNA constructs comprising ribonucleotides and 3'-terminal dithymidine caps was compared to untreated cells, scrambled siNA control constructs (Scram1 and Scram2), and cells transfected with lipid alone (transfection control). As shown in the figure, all of the siNA constructs show significant reduction of BACE RNA expression.

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Figure 31 shows a non-limiting example of reduction of cyclin D1 mRNA in A549 cells mediated by chemically-modified siNAs that target cyclin D1 mRNA. A549 cells were transfected with 0.25 ug/well of lipid complexed with 25 nM siNA. A siNA construct comprising ribonucleotides and 3'-terminal dithymidine (RPI#31009/31085) was compared to a chemically modified siNA construct comprising 2'-deoxy-2'-fluoro pyrimidine nucleotides and purine ribonucleotides in which the sense strand of the siNA is further modified with 5' and 3'-terminal inverted deoxyabasic caps and the antisense strand comprises a 3'-terminal phosphorothioate internucleotide linkage (RPI#31304/31305), which was also compared to a matched chemistry inverted control (RPI#31316/31317). In addition, the siNA constructs were also compared to untreated cells, cells transfected with lipid and scrambled siNA constructs (Scram1 and Scram2), and cells transfected with lipid alone (transfection control). As shown in the figure, both siNA constructs show significant reduction of cyclin D1 RNA expression.

Figure 32 shows a non-limiting example of reduction of PTP-1B mRNA in A549 cells mediated by chemically-modified siNAs that target PTP-1B mRNA. A549 cells were transfected with 0.25 ug/well of lipid complexed with 25 nM siNA. A siNA construct comprising ribonucleotides and 3'-terminal dithymidine caps (RPI#31018/31307) was compared to a chemically modified siNA construct comprising 2'-deoxy-2'-fluoro pyrimidine nucleotides and purine ribonucleotides in which the sense strand of the siNA is further modified with 5' and 3'-terminal inverted deoxyabasic caps and the antisense strand comprises a 3'-terminal phosphorothioate internucleotide linkage

(RPI#31306/31307), which was also compared to a matched chemistry inverted control (RPI#31318/31319). In addition, the siNA constructs were also compared to untreated cells, cells transfected with lipid and scrambled siNA constructs (Scram1 and Scram2), and cells transfected with lipid alone (transfection control). As shown in the figure, both siNA constructs show significant reduction of PTP-1B RNA expression.

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Figure 33 shows a non-limiting example of reduction of ERG2 mRNA in DLD1 cells mediated by siNAs that target ERG2 mRNA. DLD1 cells were transfected with 0.25 ug/well of lipid complexed with 25 nM siNA. A screen of siNA constructs comprising ribonucleotides and 3'-terminal dithymidine caps was compared to untreated cells, scrambled siNA control constructs (Scram1 and Scram2), and cells transfected with lipid alone (transfection control). As shown in the figure, all of the siNA constructs show significant reduction of ERG2 RNA expression.

Figure 34 shows a non-limiting example of reduction of PCNA mRNA in A549 cells mediated by chemically-modified siNAs that target PCNA mRNA. A549 cells were transfected with 0.25 ug/well of lipid complexed with 25 nM siNA. A siNA construct comprising ribonucleotides and 3'-terminal dithymidine caps (RPI#31035/31111) was compared to a chemically modified siNA construct comprising 2'-deoxy-2'-fluoro pyrimidine nucleotides and purine ribonucleotides in which the sense strand of the siNA is further modified with 5' and 3'-terminal inverted deoxyabasic caps and the antisense strand comprises a 3'-terminal phosphorothioate internucleotide (RPI#31310/31311), which was also compared to a matched chemistry inverted control (RPI#31322/31323). In addition, the siNA constructs were also compared to untreated cells, cells transfected with lipid and scrambled siNA constructs (Scram1 and Scram2), and cells transfected with lipid alone (transfection control). As shown in the figure, both siNA constructs show significant reduction of PCNA RNA expression.

DETAILED DESCRIPTION OF THE INVENTION

Mechanism of action of Nucleic Acid Molecules of the Invention

The discussion that follows discusses the proposed mechanism of RNA interference mediated by short interfering RNA as is presently known, and is not meant to be limiting and is not an admission of prior art. Applicant demonstrates herein that chemically-

modified short interfering nucleic acids possess similar or improved capacity to mediate RNAi as do siRNA molecules and are expected to possess improved stability and activity in vivo; therefore, this discussion is not meant to be limiting only to siRNA and can be applied to siNA as a whole. By "improved capacity to mediate RNAi" or "improved RNAi activity" is meant to include RNAi activity measured in vitro and/or in vivo where the RNAi activity is a reflection of both the ability of the siNA to mediate RNAi and the stability of the siNAs of the invention. In this invention, the product of these activities can be increased in vitro and/or in vivo compared to an all RNA siRNA or a siNA containing a plurality of ribonucleotides. In some cases, the activity or stability of the siNA molecule can be decreased (i.e., less than ten-fold), but the overall activity of the siNA molecule is enhanced in vitro and/or in vivo.

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RNA interference refers to the process of sequence specific post-transcriptional gene silencing in animals mediated by short interfering RNAs (siRNAs) (Fire et al., 1998, Nature, 391, 806). The corresponding process in plants is commonly referred to as posttranscriptional gene silencing or RNA silencing and is also referred to as quelling in The process of post-transcriptional gene silencing is thought to be an evolutionarily-conserved cellular defense mechanism used to prevent the expression of foreign genes which is commonly shared by diverse flora and phyla (Fire et al., 1999, Trends Genet., 15, 358). Such protection from foreign gene expression may have evolved in response to the production of double-stranded RNAs (dsRNAs) derived from viral infection or the random integration of transposon elements into a host genome via a cellular response that specifically destroys homologous single-stranded RNA or viral genomic RNA. The presence of dsRNA in cells triggers the RNAi response though a mechanism that has yet to be fully characterized. This mechanism appears to be different from the interferon response that results from dsRNA-mediated activation of protein kinase PKR and 2', 5'-oligoadenylate synthetase resulting in non-specific cleavage of mRNA by ribonuclease L.

The presence of long dsRNAs in cells stimulates the activity of a ribonuclease III enzyme referred to as Dicer. Dicer is involved in the processing of the dsRNA into short pieces of dsRNA known as short interfering RNAs (siRNAs) (Berstein *et al.*, 2001, *Nature*, 409, 363). Short interfering RNAs derived from Dicer activity are typically about 21 to about 23 nucleotides in length and comprise about 19 base pair duplexes. Dicer has

also been implicated in the excision of 21- and 22-nucleotide small temporal RNAs (stRNAs) from precursor RNA of conserved structure that are implicated in translational control (Hutvagner et al., 2001, Science, 293, 834). The RNAi response also features an endonuclease complex containing a siRNA, commonly referred to as an RNA-induced silencing complex (RISC), which mediates cleavage of single-stranded RNA having sequence homologous to the siRNA. Cleavage of the target RNA takes place in the middle of the region complementary to the guide sequence of the siRNA duplex (Elbashir et al., 2001, Genes Dev., 15, 188). In addition, RNA interference can also involve small RNA (e.g., micro-RNA or miRNA) mediated gene silencing, presumably though cellular mechanisms that regulate chromatin structure and thereby prevent transcription of target gene sequences (see for example Allshire, 2002, Science, 297, 1818-1819; Volpe et al., 2002, Science, 297, 1833-1837; Jenuwein, 2002, Science, 297, 2215-2218; and Hall et al., 2002, Science, 297, 2232-2237). As such, siNA molecules of the invention can be used to mediate gene silencing via interaction with RNA transcripts or alternately by interaction with particular gene sequences, wherein such interaction results in gene silencing either at the transcriptional level or post-transcriptional level.

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RNAi has been studied in a variety of systems. Fire et al., 1998, Nature, 391, 806, were the first to observe RNAi in C. elegans. Wianny and Goetz, 1999, Nature Cell Biol., 2, 70, describe RNAi mediated by dsRNA in mouse embryos. Hammond et al., 2000, Nature, 404, 293, describe RNAi in Drosophila cells transfected with dsRNA. Elbashir et al., 2001, Nature, 411, 494, describe RNAi induced by introduction of duplexes of synthetic 21-nucleotide RNAs in cultured mammalian cells including human embryonic kidney and HeLa cells. Recent work in Drosophila embryonic lysates has revealed certain requirements for siRNA length, structure, chemical composition, and sequence that are essential to mediate efficient RNAi activity. These studies have shown that 21 nucleotide siRNA duplexes are most active when containing two 2-nucleotide 3'terminal nucleotide overhangs. Furthermore, substitution of one or both siRNA strands with 2'-deoxy or 2'-O-methyl nucleotides abolishes RNAi activity, whereas substitution of 3'-terminal siRNA nucleotides with deoxy nucleotides was shown to be tolerated. Mismatch sequences in the center of the siRNA duplex were also shown to abolish RNAi activity. In addition, these studies also indicate that the position of the cleavage site in the target RNA is defined by the 5'-end of the siRNA guide sequence rather than the 3'-end

(Elbashir et al., 2001, EMBO J., 20, 6877). Other studies have indicated that a 5'-phosphate on the target-complementary strand of a siRNA duplex is required for siRNA activity and that ATP is utilized to maintain the 5'-phosphate moiety on the siRNA (Nykanen et al., 2001, Cell, 107, 309); however, siRNA molecules lacking a 5'-phosphate are active when introduced exogenously, suggesting that 5'-phosphorylation of siRNA constructs may occur in vivo.

Synthesis of Nucleic acid Molecules

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Synthesis of nucleic acids greater than 100 nucleotides in length is difficult using automated methods, and the therapeutic cost of such molecules is prohibitive. In this invention, small nucleic acid motifs "small" refers to nucleic acid motifs no more than 100 nucleotides in length, preferably no more than 80 nucleotides in length, and most preferably no more than 50 nucleotides in length; e.g., individual siNA oligonucleotide sequences or siNA sequences synthesized in tandem) are preferably used for exogenous delivery. The simple structure of these molecules increases the ability of the nucleic acid to invade targeted regions of protein and/or RNA structure. Exemplary molecules of the instant invention are chemically synthesized, and others can similarly be synthesized.

Oligonucleotides (e.g., certain modified oligonucleotides or portions of oligonucleotides lacking ribonucleotides) are synthesized using protocols known in the art, for example as described in Caruthers et al., 1992, Methods in Enzymology 211, 3-19, Thompson et al., International PCT Publication No. WO 99/54459, Wincott et al., 1995, Nucleic Acids Res. 23, 2677-2684, Wincott et al., 1997, Methods Mol. Bio., 74, 59, Brennan et al., 1998, Biotechnol Bioeng., 61, 33-45, and Brennan, U.S. Pat. No. 6,001,311. All of these references are incorporated herein by reference. The synthesis of oligonucleotides makes use of common nucleic acid protecting and coupling groups, such as dimethoxytrityl at the 5'-end, and phosphoramidites at the 3'-end. In a non-limiting example, small scale syntheses are conducted on a 394 Applied Biosystems, Inc. synthesizer using a 0.2 µmol scale protocol with a 2.5 min coupling step for 2'-deoxy-2'-fluoro nucleotides. Table II outlines the amounts and the contact times of the reagents used in the synthesis cycle. Alternatively, syntheses at the 0.2 µmol scale can be performed on a 96-well plate synthesizer, such as the instrument produced by Protogene

(Palo Alto, CA) with minimal modification to the cycle. A 33-fold excess (60 µL of 0.11 M = 6.6 μmol) of 2'-O-methyl phosphoramidite and a 105-fold excess of S-ethyl tetrazole (60 μ L of 0.25 M = 15 μ mol) can be used in each coupling cycle of 2'-O-methyl residues relative to polymer-bound 5'-hydroxyl. A 22-fold excess (40 μ L of 0.11 M = 4.4 μ mol) of deoxy phosphoramidite and a 70-fold excess of S-ethyl tetrazole (40 μ L of 0.25 M = 10 µmol) can be used in each coupling cycle of deoxy residues relative to polymer-bound 5'-hydroxyl. Average coupling yields on the 394 Applied Biosystems, Inc. synthesizer, determined by colorimetric quantitation of the trityl fractions, are typically 97.5-99%. Other oligonucleotide synthesis reagents for the 394 Applied Biosystems, Inc. synthesizer include the following: detritylation solution is 3% TCA in methylene chloride (ABI); capping is performed with 16% N-methyl imidazole in THF (ABI) and 10% acetic anhydride/10% 2,6-lutidine in THF (ABI); and oxidation solution is 16.9 mM I₂, 49 mM pyridine, 9% water in THF (PERSEPTIVE™). Burdick & Jackson Synthesis Grade acetonitrile is used directly from the reagent bottle. S-Ethyltetrazole solution (0.25 M in acetonitrile) is made up from the solid obtained from American International Chemical, Inc. Alternately, for the introduction of phosphorothioate linkages, Beaucage reagent (3H-1,2-Benzodithiol-3-one 1,1-dioxide, 0.05 M in acetonitrile) is used.

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Deprotection of the DNA-based oligonucleotides is performed as follows: the polymer-bound trityl-on oligoribonucleotide is transferred to a 4 mL glass screw top vial and suspended in a solution of 40% aq. methylamine (1 mL) at 65 °C for 10 min. After cooling to -20 °C, the supernatant is removed from the polymer support. The support is washed three times with 1.0 mL of EtOH:MeCN:H2O/3:1:1, vortexed and the supernatant is then added to the first supernatant. The combined supernatants, containing the oligoribonucleotide, are dried to a white powder.

The method of synthesis used for RNA including certain siNA molecules of the invention follows the procedure as described in Usman et al., 1987, J. Am. Chem. Soc., 109, 7845; Scaringe et al., 1990, Nucleic Acids Res., 18, 5433; and Wincott et al., 1995, Nucleic Acids Res. 23, 2677-2684 Wincott et al., 1997, Methods Mol. Bio., 74, 59, and makes use of common nucleic acid protecting and coupling groups, such as dimethoxytrityl at the 5'-end, and phosphoramidites at the 3'-end. In a non-limiting example, small scale syntheses are conducted on a 394 Applied Biosystems, Inc. synthesizer using a 0.2 µmol scale protocol with a 7.5 min coupling step for alkylsilyl

protected nucleotides and a 2.5 min coupling step for 2'-O-methylated nucleotides. Table II outlines the amounts and the contact times of the reagents used in the synthesis cycle. Alternatively, syntheses at the 0.2 µmol scale can be done on a 96-well plate synthesizer, such as the instrument produced by Protogene (Palo Alto, CA) with minimal modification to the cycle. A 33-fold excess (60 μ L of 0.11 M = 6.6 μ mol) of 2'-O-methyl phosphoramidite and a 75-fold excess of S-ethyl tetrazole (60 μ L of 0.25 M = 15 μ mol) can be used in each coupling cycle of 2'-O-methyl residues relative to polymer-bound 5'hydroxyl. A 66-fold excess (120 µL of 0.11 M = 13.2 µmol) of alkylsilyl (ribo) protected phosphoramidite and a 150-fold excess of S-ethyl tetrazole (120 μ L of 0.25 M = 30 μ mol) can be used in each coupling cycle of ribo residues relative to polymer-bound 5'hydroxyl. Average coupling yields on the 394 Applied Biosystems, Inc. synthesizer, determined by colorimetric quantitation of the trityl fractions, are typically 97.5-99%. Other oligonucleotide synthesis reagents for the 394 Applied Biosystems, Inc. synthesizer include the following: detritylation solution is 3% TCA in methylene chloride (ABI); capping is performed with 16% N-methyl imidazole in THF (ABI) and 10% acetic anhydride/10% 2,6-lutidine in THF (ABI); oxidation solution is 16.9 mM I2, 49 mM pyridine, 9% water in THF (PERSEPTIVE™). Burdick & Jackson Synthesis Grade acetonitrile is used directly from the reagent bottle. S-Ethyltetrazole solution (0.25 M in acetonitrile) is made up from the solid obtained from American International Chemical, Inc. Alternately, for the introduction of phosphorothioate linkages, Beaucage reagent (3H-1,2-Benzodithiol-3-one 1,1-dioxide0.05 M in acetonitrile) is used.

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Deprotection of the RNA is performed using either a two-pot or one-pot protocol. For the two-pot protocol, the polymer-bound trityl-on oligoribonucleotide is transferred to a 4 mL glass screw top vial and suspended in a solution of 40% aq. methylamine (1 mL) at 65 °C for 10 min. After cooling to -20 °C, the supernatant is removed from the polymer support. The support is washed three times with 1.0 mL of EtOH:MeCN:H2O/3:1:1, vortexed and the supernatant is then added to the first supernatant. The combined supernatants, containing the oligoribonucleotide, are dried to a white powder. The base deprotected oligoribonucleotide is resuspended in anhydrous TEA/HF/NMP solution (300 μL of a solution of 1.5 mL N-methylpyrrolidinone, 750 μL TEA and 1 mL TEA•3HF to provide a 1.4 M HF concentration) and heated to 65 °C. After 1.5 h, the oligomer is quenched with 1.5 M NH₄HCO₃.

Alternatively, for the one-pot protocol, the polymer-bound trityl-on oligoribonucleotide is transferred to a 4 mL glass screw top vial and suspended in a solution of 33% ethanolic methylamine/DMSO: 1/1 (0.8 mL) at 65 °C for 15 min. The vial is brought to rt. TEA•3HF (0.1 mL) is added and the vial is heated at 65 °C for 15 min. The sample is cooled at -20 °C and then quenched with 1.5 M NH₄HCO₃.

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For purification of the trityl-on oligomers, the quenched NH₄HCO₃ solution is loaded onto a C-18 containing cartridge that had been prewashed with acetonitrile followed by 50 mM TEAA. After washing the loaded cartridge with water, the RNA is detritylated with 0.5% TFA for 13 min. The cartridge is then washed again with water, salt exchanged with 1 M NaCl and washed with water again. The oligonucleotide is then eluted with 30% acetonitrile.

The average stepwise coupling yields are typically >98% (Wincott et al., 1995 Nucleic Acids Res. 23, 2677-2684). Those of ordinary skill in the art will recognize that the scale of synthesis can be adapted to be larger or smaller than the example described above including but not limited to 96-well format.

Alternatively, the nucleic acid molecules of the present invention can be synthesized separately and joined together post-synthetically, for example, by ligation (Moore et al., 1992, Science 256, 9923; Draper et al., International PCT publication No. WO 93/23569; Shabarova et al., 1991, Nucleic Acids Research 19, 4247; Bellon et al., 1997, Nucleosides & Nucleotides, 16, 951; Bellon et al., 1997, Bioconjugate Chem. 8, 204), or by hybridization following synthesis and/or deprotection.

The siNA molecules of the invention can also be synthesized via a tandem synthesis methodology as described in Example 1 herein, wherein both siNA strands are synthesized as a single contiguous oligonucleotide fragment or strand separated by a cleavable linker which is subsequently cleaved to provide separate siNA fragments or strands that hybridize and permit purification of the siNA duplex. The linker can be a polynucleotide linker or a non-nucleotide linker. The tandem synthesis of siNA as described herein can be readily adapted to both multiwell/multiplate synthesis platforms such as 96 well or similarly larger multi-well platforms. The tandem synthesis of siNA as

described herein can also be readily adapted to large scale synthesis platforms employing batch reactors, synthesis columns and the like.

A siNA molecule can also be assembled from two distinct nucleic acid strands or fragments wherein one fragment includes the sense region and the second fragment includes the antisense region of the RNA molecule.

The nucleic acid molecules of the present invention can be modified extensively to enhance stability by modification with nuclease resistant groups, for example, 2'-amino, 2'-C-allyl, 2'-fluoro, 2'-O-methyl, 2'-H (for a review see Usman and Cedergren, 1992, TIBS 17, 34; Usman et al., 1994, Nucleic Acids Symp. Ser. 31, 163). siNA constructs can be purified by gel electrophoresis using general methods or can be purified by high pressure liquid chromatography (HPLC; see Wincott et al., supra, the totality of which is hereby incorporated herein by reference) and re-suspended in water.

In another aspect of the invention, siNA molecules of the invention are expressed from transcription units inserted into DNA or RNA vectors. The recombinant vectors can be DNA plasmids or viral vectors. siNA expressing viral vectors can be constructed based on, but not limited to, adeno-associated virus, retrovirus, adenovirus, or alphavirus. The recombinant vectors capable of expressing the siNA molecules can be delivered as described herein, and persist in target cells. Alternatively, viral vectors can be used that provide for transient expression of siNA molecules.

20 Optimizing Activity of the nucleic acid molecule of the invention.

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Chemically synthesizing nucleic acid molecules with modifications (base, sugar and/or phosphate) can prevent their degradation by serum ribonucleases, which can increase their potency (see e.g., Eckstein et al., International Publication No. WO 92/07065; Perrault et al., 1990 Nature 344, 565; Pieken et al., 1991, Science 253, 314; Usman and Cedergren, 1992, Trends in Biochem. Sci. 17, 334; Usman et al., International Publication No. WO 93/15187; and Rossi et al., International Publication No. WO 91/03162; Sproat, U.S. Pat. No. 5,334,711; Gold et al., U.S. Pat. No. 6,300,074; and Burgin et al., supra; all of which are incorporated by reference herein). All of the above references describe various chemical modifications that can be made to the base, phosphate and/or sugar moieties of the nucleic acid molecules described herein.

Modifications that enhance their efficacy in cells, and removal of bases from nucleic acid molecules to shorten oligonucleotide synthesis times and reduce chemical requirements are desired.

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There are several examples in the art describing sugar, base and phosphate modifications that can be introduced into nucleic acid molecules with significant enhancement in their nuclease stability and efficacy. For example, oligonucleotides are modified to enhance stability and/or enhance biological activity by modification with nuclease resistant groups, for example, 2'-amino, 2'-C-allyl, 2'-fluoro, 2'-O-methyl, 2'-Oallyl, 2'-H, nucleotide base modifications (for a review see Usman and Cedergren, 1992, TIBS. 17, 34; Usman et al., 1994, Nucleic Acids Symp. Ser. 31, 163; Burgin et al., 1996, Biochemistry, 35, 14090). Sugar modification of nucleic acid molecules have been extensively described in the art (see Eckstein et al., International Publication PCT No. WO 92/07065; Perrault et al. Nature, 1990, 344, 565-568; Pieken et al. Science, 1991, 253, 314-317; Usman and Cedergren, Trends in Biochem. Sci., 1992, 17, 334-339; Usman et al. International Publication PCT No. WO 93/15187; Sproat, U.S. Pat. No. 5,334,711 and Beigelman et al., 1995, J. Biol. Chem., 270, 25702; Beigelman et al., International PCT publication No. WO 97/26270; Beigelman et al., U.S. Pat. No. 5.716.824; Usman et al., U.S. Pat. No. 5,627,053; Woolf et al., International PCT Publication No. WO 98/13526; Thompson et al., USSN 60/082,404 which was filed on April 20, 1998; Karpeisky et al., 1998, Tetrahedron Lett., 39, 1131; Earnshaw and Gait, 1998, Biopolymers (Nucleic Acid Sciences), 48, 39-55; Verma and Eckstein, 1998, Annu. Rev. Biochem., 67, 99-134; and Burlina et al., 1997, Bioorg. Med. Chem., 5, 1999-2010; all of the references are hereby incorporated in their totality by reference herein). Such publications describe general methods and strategies to determine the location of incorporation of sugar, base and/or phosphate modifications and the like into nucleic acid molecules without modulating catalysis, and are incorporated by reference herein. In view of such teachings, similar modifications can be used as described herein to modify the siNA nucleic acid molecules of the instant invention so long as the ability of siNA to promote RNAi is cells is not significantly inhibited.

While chemical modification of oligonucleotide internucleotide linkages with phosphorothioate, phosphorodithioate, and/or 5'-methylphosphonate linkages improves stability, excessive modifications can cause some toxicity or decreased activity.

Therefore, when designing nucleic acid molecules, the amount of these internucleotide linkages should be minimized. The reduction in the concentration of these linkages should lower toxicity, resulting in increased efficacy and higher specificity of these molecules.

Short interfering nucleic acid (siNA) molecules having chemical modifications that maintain or enhance activity are provided. Such a nucleic acid is also generally more resistant to nucleases than an unmodified nucleic acid. Accordingly, the *in vitro* and/or *in vivo* activity should not be significantly lowered. In cases in which modulation is the goal, therapeutic nucleic acid molecules delivered exogenously should optimally be stable within cells until translation of the target RNA has been modulated long enough to reduce the levels of the undesirable protein. This period of time varies between hours to days depending upon the disease state. Improvements in the chemical synthesis of RNA and DNA (Wincott *et al.*, 1995, *Nucleic Acids Res.* 23, 2677; Caruthers *et al.*, 1992, *Methods in Enzymology* 211,3-19 (incorporated by reference herein)) have expanded the ability to modify nucleic acid molecules by introducing nucleotide modifications to enhance their nuclease stability, as described above.

In one embodiment, nucleic acid molecules of the invention include one or more (e.g., about 1, 2, 3, 4, 5, 6, 7, 8, 9, 10, or more) G-clamp nucleotides. A G-clamp nucleotide is a modified cytosine analog wherein the modifications confer the ability to hydrogen bond both Watson-Crick and Hoogsteen faces of a complementary guanine within a duplex, see for example Lin and Matteucci, 1998, J. Am. Chem. Soc., 120, 8531-8532. A single G-clamp analog substitution within an oligonucleotide can result in substantially enhanced helical thermal stability and mismatch discrimination when hybridized to complementary oligonucleotides. The inclusion of such nucleotides in nucleic acid molecules of the invention results in both enhanced affinity and specificity to nucleic acid targets, complementary sequences, or template strands. In another embodiment, nucleic acid molecules of the invention include one or more (e.g., about 1, 2, 3, 4, 5, 6, 7, 8, 9, 10, or more) LNA "locked nucleic acid" nucleotides such as a 2', 4'-C methylene bicyclo nucleotide (see for example Wengel et al., International PCT Publication No. WO 00/66604 and WO 99/14226).

In another embodiment, the invention features conjugates and/or complexes of siNA molecules of the invention. Such conjugates and/or complexes can be used to facilitate delivery of siNA molecules into a biological system, such as a cell. The conjugates and complexes provided by the instant invention can impart therapeutic activity by transferring therapeutic compounds across cellular membranes, altering the pharmacokinetics, and/or modulating the localization of nucleic acid molecules of the The present invention encompasses the design and synthesis of novel invention. conjugates and complexes for the delivery of molecules, including, but not limited to, small molecules, lipids, phospholipids, nucleosides, nucleotides, nucleic acids, antibodies, toxins, negatively charged polymers and other polymers, for example proteins, peptides, hormones, carbohydrates, polyethylene glycols, or polyamines, across cellular membranes. In general, the transporters described are designed to be used either individually or as part of a multi-component system, with or without degradable linkers. These compounds are expected to improve delivery and/or localization of nucleic acid molecules of the invention into a number of cell types originating from different tissues. in the presence or absence of serum (see Sullenger and Cech, U.S. Pat. No. 5,854,038). Conjugates of the molecules described herein can be attached to biologically active molecules via linkers that are biodegradable, such as biodegradable nucleic acid linker molecules.

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The term "biodegradable linker" as used herein, refers to a nucleic acid or non-nucleic acid linker molecule that is designed as a biodegradable linker to connect one molecule to another molecule, for example, a biologically active molecule to a siNA molecule of the invention or the sense and antisense strands of a siNA molecule of the invention. The biodegradable linker is designed such that its stability can be modulated for a particular purpose, such as delivery to a particular tissue or cell type. The stability of a nucleic acid-based biodegradable linker molecule can be modulated by using various chemistries, for example combinations of ribonucleotides, deoxyribonucleotides, and chemically-modified nucleotides, such as 2'-O-methyl, 2'-fluoro, 2'-amino, 2'-O-amino, 2'-C-allyl, 2'-O-allyl, and other 2'-modified or base modified nucleotides. The biodegradable nucleic acid linker molecule can be a dimer, trimer, tetramer or longer nucleic acid molecule, for example, an oligonucleotide of about 2, 3, 4, 5, 6, 7, 8, 9, 10, 11, 12, 13, 14, 15, 16, 17, 18, 19, or 20 nucleotides in length, or can comprise a single

nucleotide with a phosphorus-based linkage, for example, a phosphoramidate or phosphodiester linkage. The biodegradable nucleic acid linker molecule can also comprise nucleic acid backbone, nucleic acid sugar, or nucleic acid base modifications.

The term "biodegradable" as used herein, refers to degradation in a biological system, for example enzymatic degradation or chemical degradation.

The term "biologically active molecule" as used herein, refers to compounds or molecules that are capable of eliciting or modifying a biological response in a system. Non-limiting examples of biologically active siNA molecules either alone or in combination with other molecules contemplated by the instant invention include therapeutically active molecules such as antibodies, hormones, antivirals, peptides, proteins, chemotherapeutics, small molecules, vitamins, co-factors, nucleosides, nucleotides, oligonucleotides, enzymatic nucleic acids, antisense nucleic acids, triplex forming oligonucleotides, 2,5-A chimeras, siNA, dsRNA, allozymes, aptamers, decoys and analogs thereof. Biologically active molecules of the invention also include molecules capable of modulating the pharmacokinetics and/or pharmacodynamics of other biologically active molecules, for example, lipids and polymers such as polyamines, polyamides, polyethylene glycol and other polyethers.

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The term "phospholipid" as used herein, refers to a hydrophobic molecule comprising at least one phosphorus group. For example, a phospholipid can comprise a phosphorus-containing group and saturated or unsaturated alkyl group, optionally substituted with OH, COOH, oxo, amine, or substituted or unsubstituted aryl groups.

Therapeutic nucleic acid molecules (e.g., siNA molecules) delivered exogenously optimally are stable within cells until reverse transcription of the RNA has been modulated long enough to reduce the levels of the RNA transcript. The nucleic acid molecules are resistant to nucleases in order to function as effective intracellular therapeutic agents. Improvements in the chemical synthesis of nucleic acid molecules described in the instant invention and in the art have expanded the ability to modify nucleic acid molecules by introducing nucleotide modifications to enhance their nuclease stability as described above.

In yet another embodiment, siNA molecules having chemical modifications that maintain or enhance enzymatic activity of proteins involved in RNAi are provided. Such nucleic acids are also generally more resistant to nucleases than unmodified nucleic acids. Thus, in vitro and/or in vivo the activity should not be significantly lowered.

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Use of the nucleic acid-based molecules of the invention will lead to better treatment of the disease progression by affording the possibility of combination therapies (e.g., multiple siNA molecules targeted to different genes; nucleic acid molecules coupled with known small molecule modulators; or intermittent treatment with combinations of molecules, including different motifs and/or other chemical or biological molecules). The treatment of subjects with siNA molecules can also include combinations of different types of nucleic acid molecules, such as enzymatic nucleic acid molecules (ribozymes), allozymes, antisense, 2,5-A oligoadenylate, decoys, and aptamers.

In another aspect a siNA molecule of the invention comprises one or more 5' and/or a 3'- cap structure, for example on only the sense siNA strand, the antisense siNA strand, or both siNA strands.

By "cap structure" is meant chemical modifications, which have been incorporated at either terminus of the oligonucleotide (see, for example, Adamic et al., U.S. Pat. No. 5,998,203, incorporated by reference herein). These terminal modifications protect the nucleic acid molecule from exonuclease degradation, and may help in delivery and/or localization within a cell. The cap may be present at the 5'-terminus (5'-cap) or at the 3'terminal (3'-cap) or may be present on both termini. In non-limiting examples, the 5'-cap is selected from the group consisting of glyceryl, inverted deoxy abasic residue (moiety); 4',5'-methylene nucleotide; 1-(beta-D-erythrofuranosyl) nucleotide, 4'-thio nucleotide; carbocyclic nucleotide; 1,5-anhydrohexitol nucleotide; L-nucleotides; alpha-nucleotides; modified base nucleotide; phosphorodithioate linkage; threo-pentofuranosyl nucleotide; acyclic 3',4'-seco nucleotide; acyclic 3,4-dihydroxybutyl nucleotide; acyclic 3,5dihydroxypentyl nucleotide, 3'-3'-inverted nucleotide moiety; 3'-3'-inverted abasic moiety; 3'-2'-inverted nucleotide moiety; 3'-2'-inverted abasic moiety; 1,4-butanediol phosphate; 3'-phosphoramidate; hexylphosphate; aminohexyl phosphate; 3'-phosphate; 3'phosphorothioate; phosphorodithioate; or bridging or non-bridging methylphosphonate moiety.

In non-limiting examples, the 3'-cap is selected from the group consisting of glyceryl, inverted deoxy abasic residue (moiety), 4',5'-methylene nucleotide; 1-(beta-Derythrofuranosyl) nucleotide; 4'-thio nucleotide, carbocyclic nucleotide; 5'-amino-alkyl phosphate; 1,3-diamino-2-propyl phosphate; 3-aminopropyl phosphate; 6-aminohexyl phosphate; 1,2-aminododecyl phosphate; hydroxypropyl phosphate; 1,5-anhydrohexitol nucleotide; L-nucleotide; alpha-nucleotide; modified base nucleotide; phosphorodithioate; threo-pentofuranosyl nucleotide; acyclic 3',4'-seco nucleotide; 3,4-dihydroxybutyl nucleotide; 3,5-dihydroxypentyl nucleotide, 5'-5'-inverted nucleotide moiety; 5'-5'-inverted abasic moiety; 5'-phosphoramidate; 5'-phosphorothioate; 1,4-butanediol phosphate; 5'-amino; bridging and/or non-bridging 5'-phosphoramidate, phosphorothioate and/or phosphorodithioate, bridging or non bridging methylphosphonate and 5'-mercapto moieties (for more details see Beaucage and Iyer, 1993, Tetrahedron 49, 1925; incorporated by reference herein).

By the term "non-nucleotide" is meant any group or compound which can be incorporated into a nucleic acid chain in the place of one or more nucleotide units, including either sugar and/or phosphate substitutions, and allows the remaining bases to exhibit their enzymatic activity. The group or compound is abasic in that it does not contain a commonly recognized nucleotide base, such as adenosine, guanine, cytosine, uracil or thymine and therefore lacks a base at the 1'-position.

An "alkyl" group refers to a saturated aliphatic hydrocarbon, including straight-chain, branched-chain, and cyclic alkyl groups. Preferably, the alkyl group has 1 to 12 carbons. More preferably, it is a lower alkyl of from 1 to 7 carbons, more preferably 1 to 4 carbons. The alkyl group can be substituted or unsubstituted. When substituted the substituted group(s) is preferably, hydroxyl, cyano, alkoxy, =O, =S, NO2 or N(CH3)2, amino, or SH. The term also includes alkenyl groups that are unsaturated hydrocarbon groups containing at least one carbon-carbon double bond, including straight-chain, branched-chain, and cyclic groups. Preferably, the alkenyl group has 1 to 12 carbons. More preferably, it is a lower alkenyl of from 1 to 7 carbons, more preferably 1 to 4 carbons. The alkenyl group may be substituted or unsubstituted. When substituted the substituted group(s) is preferably, hydroxyl, cyano, alkoxy, =O, =S, NO2, halogen, N(CH3)2, amino, or SH. The term "alkyl" also includes alkynyl groups that have an

unsaturated hydrocarbon group containing at least one carbon-carbon triple bond, including straight-chain, branched-chain, and cyclic groups. Preferably, the alkynyl group has 1 to 12 carbons. More preferably, it is a lower alkynyl of from 1 to 7 carbons, more preferably 1 to 4 carbons. The alkynyl group may be substituted or unsubstituted. When substituted the substituted group(s) is preferably, hydroxyl, cyano, alkoxy, =O, =S, NO2 or N(CH₃)₂, amino or SH.

Such alkyl groups can also include aryl, alkylaryl, carbocyclic aryl, heterocyclic aryl, amide and ester groups. An "aryl" group refers to an aromatic group that has at least one ring having a conjugated pi electron system and includes carbocyclic aryl, heterocyclic aryl and biaryl groups, all of which may be optionally substituted. The preferred substituent(s) of aryl groups are halogen, trihalomethyl, hydroxyl, SH, OH, cyano, alkoxy, alkyl, alkenyl, alkynyl, and amino groups. An "alkylaryl" group refers to an alkyl group (as described above) covalently joined to an aryl group (as described above). Carbocyclic aryl groups are groups wherein the ring atoms on the aromatic ring are all carbon atoms. The carbon atoms are optionally substituted. Heterocyclic aryl groups are groups having from 1 to 3 heteroatoms as ring atoms in the aromatic ring and the remainder of the ring atoms are carbon atoms. Suitable heteroatoms include oxygen, sulfur, and nitrogen, and include furanyl, thienyl, pyridyl, pyrrolyl, N-lower alkyl pyrrolo, pyrimidyl, pyrazinyl, imidazolyl and the like, all optionally substituted. An "amide" refers to an -C(O)-NH-R, where R is either alkyl, aryl, alkylaryl or hydrogen.

By "nucleotide" as used herein is as recognized in the art to include natural bases (standard), and modified bases well known in the art. Such bases are generally located at the 1' position of a nucleotide sugar moiety. Nucleotides generally comprise a base, sugar and a phosphate group. The nucleotides can be unmodified or modified at the sugar, phosphate and/or base moiety, (also referred to interchangeably as nucleotide analogs, modified nucleotides, non-natural nucleotides, non-standard nucleotides and other; see, for example, Usman and McSwiggen, supra; Eckstein et al., International PCT Publication No. WO 92/07065; Usman et al., International PCT Publication No. WO 93/15187; Uhlman & Peyman, supra, all are hereby incorporated by reference herein). There are several examples of modified nucleic acid bases known in the art as summarized by Limbach et al., 1994, Nucleic Acids Res. 22, 2183. Some of the non-

limiting examples of base modifications that can be introduced into nucleic acid molecules include, inosine, purine, pyridin-4-one, pyridin-2-one, phenyl, pseudouracil, 2, 4, 6-trimethoxy benzene, 3-methyl uracil, dihydrouridine, naphthyl, aminophenyl, 5-alkylcytidines (e.g., 5-methylcytidine), 5-alkyluridines (e.g., ribothymidine), 5-halouridine (e.g., 5-bromouridine) or 6-azapyrimidines or 6-alkylpyrimidines (e.g. 6-methyluridine), propyne, and others (Burgin et al., 1996, Biochemistry, 35, 14090; Uhlman & Peyman, supra). By "modified bases" in this aspect is meant nucleotide bases other than adenine, guanine, cytosine and uracil at 1' position or their equivalents.

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In one embodiment, the invention features modified siNA molecules, with phosphate backbone modifications comprising one or more phosphorothioate, phosphorodithioate, methylphosphonate, phosphotriester, morpholino, amidate carbamate, carboxymethyl, acetamidate, polyamide, sulfonate, sulfonamide, sulfamate, formacetal, thioformacetal, and/or alkylsilyl, substitutions. For a review of oligonucleotide backbone modifications, see Hunziker and Leumann, 1995, Nucleic Acid Analogues: Synthesis and Properties, in Modern Synthetic Methods, VCH, 331-417, and Mesmaeker et al., 1994, Novel Backbone Replacements for Oligonucleotides, in Carbohydrate Modifications in Antisense Research, ACS, 24-39.

By "abasic" is meant sugar moieties lacking a base or having other chemical groups in place of a base at the 1' position, see for example Adamic *et al.*, U.S. Pat. No. 5,998,203.

By "unmodified nucleoside" is meant one of the bases adenine, cytosine, guanine, thymine, or uracil joined to the 1' carbon of β -D-ribo-furanose.

By "modified nucleoside" is meant any nucleotide base which contains a modification in the chemical structure of an unmodified nucleotide base, sugar and/or phosphate. Non-limiting examples of modified nucleotides are shown by Formulae I-VII and/or other modifications described herein.

In connection with 2'-modified nucleotides as described for the present invention, by "amino" is meant 2'-NH₂ or 2'-O-NH₂, which can be modified or unmodified. Such modified groups are described, for example, in Eckstein *et al.*, U.S. Pat. No. 5,672,695

and Matulic-Adamic et al., U.S. Pat. No. 6,248,878, which are both incorporated by reference in their entireties.

Various modifications to nucleic acid siNA structure can be made to enhance the utility of these molecules. Such modifications will enhance shelf-life, half-life in vitro, stability, and ease of introduction of such oligonucleotides to the target site, e.g., to enhance penetration of cellular membranes, and confer the ability to recognize and bind to targeted cells.

Administration of Nucleic Acid Molecules

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A siNA molecule of the invention can be adapted for use to treat any disease, infection or condition associated with gene expression, and other indications that can respond to the level of gene product in a cell or tissue, alone or in combination with other therapies. For example, a siNA molecule can comprise a delivery vehicle, including liposomes, for administration to a subject, carriers and diluents and their salts, and/or can be present in pharmaceutically acceptable formulations. Methods for the delivery of nucleic acid molecules are described in Akhtar et al., 1992, Trends Cell Bio., 2, 139; Delivery Strategies for Antisense Oligonucleotide Therapeutics, ed. Akhtar, 1995, Maurer et al., 1999, Mol. Membr. Biol., 16, 129-140; Hofland and Huang, 1999, Handb. Exp. Pharmacol., 137, 165-192; and Lee et al., 2000, ACS Symp. Ser., 752, 184-192, all of which are incorporated herein by reference. Beigelman et al., U.S. Pat. No. 6,395,713 and Sullivan et al., PCT WO 94/02595 further describe the general methods for delivery of nucleic acid molecules. These protocols can be utilized for the delivery of virtually any nucleic acid molecule. Nucleic acid molecules can be administered to cells by a variety of methods known to those of skill in the art, including, but not restricted to, encapsulation in liposomes, by iontophoresis, or by incorporation into other vehicles, such as hydrogels, cyclodextrins (see for example Gonzalez et al., 1999, Bioconjugate Chem., 10, 1068-1074), biodegradable nanocapsules, and bioadhesive microspheres, or by proteinaceous vectors (O'Hare and Normand, International PCT Publication No. WO 00/53722). Alternatively, the nucleic acid/vehicle combination is locally delivered by direct injection or by use of an infusion pump. Direct injection of the nucleic acid molecules of the invention, whether subcutaneous, intramuscular, or intradermal, can take place using standard needle and syringe methodologies, or by needle-free technologies

such as those described in Conry et al., 1999, Clin. Cancer Res., 5, 2330-2337 and Barry et al., International PCT Publication No. WO 99/31262. Many examples in the art describe CNS delivery methods of oligonucleotides by osmotic pump, (see Chun et al., 1998, Neuroscience Letters, 257, 135-138, D'Aldin et al., 1998, Mol. Brain Research, 55, 151-164, Dryden et al., 1998, J. Endocrinol., 157, 169-175, Ghirnikar et al., 1998, Neuroscience Letters, 247, 21-24) or direct infusion (Broaddus et al., 1997, Neurosurg. Focus, 3, article 4). Other routes of delivery include, but are not limited to oral (tablet or pill form) and/or intrathecal delivery (Gold, 1997, Neuroscience, 76, 1153-1158). More detailed descriptions of nucleic acid delivery and administration are provided in Sullivan et al., supra, Draper et al., PCT WO93/23569, Beigelman et al., PCT WO99/05094, and Klimuk et al., PCT WO99/04819 all of which have been incorporated by reference herein. The molecules of the instant invention can be used as pharmaceutical agents. Pharmaceutical agents prevent, modulate the occurrence, or treat (alleviate a symptom to some extent, preferably all of the symptoms) of a disease state in a subject.

In addition, the invention features the use of methods to deliver the nucleic acid molecules of the instant invention to hematopoietic cells, including monocytes and lymphocytes. These methods are described in detail by Hartmann *et al.*, 1998, *J. Phamacol. Exp. Ther.*, 285(2), 920-928; Kronenwett *et al.*, 1998, *Blood*, 91(3), 852-862; Filion and Phillips, 1997, *Biochim. Biophys. Acta.*, 1329(2), 345-356; Ma and Wei, 1996, *Leuk. Res.*, 20(11/12), 925-930; and Bongartz *et al.*, 1994, *Nucleic Acids Research*, 22(22), 4681-8. Such methods, as described above, include the use of free oligonucleitide, cationic lipid formulations, liposome formulations including pH sensitive liposomes and immunoliposomes, and bioconjugates including oligonucleotides conjugated to fusogenic peptides, for the transfection of hematopoietic cells with oligonucleotides.

Thus, the invention features a pharmaceutical composition comprising one or more nucleic acid(s) of the invention in an acceptable carrier, such as a stabilizer, buffer, and the like. The polynucleotides of the invention can be administered (e.g., RNA, DNA or protein) and introduced into a subject by any standard means, with or without stabilizers, buffers, and the like, to form a pharmaceutical composition. When it is desired to use a liposome delivery mechanism, standard protocols for formation of liposomes can be followed. The compositions of the present invention can also be formulated and used as

tablets, capsules or elixirs for oral administration, suppositories for rectal administration, sterile solutions, suspensions for injectable administration, and the other compositions known in the art.

The present invention also includes pharmaceutically acceptable formulations of the compounds described. These formulations include salts of the above compounds, e.g., acid addition salts, for example, salts of hydrochloric, hydrobromic, acetic acid, and benzene sulfonic acid.

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A pharmacological composition or formulation refers to a composition or formulation in a form suitable for administration, e.g., systemic administration, into a cell or subject, including for example a human. Suitable forms, in part, depend upon the use or the route of entry, for example oral, transdermal, or by injection. Such forms should not prevent the composition or formulation from reaching a target cell (i.e., a cell to which the negatively charged nucleic acid is desirable for delivery). For example, pharmacological compositions injected into the blood stream should be soluble. Other factors are known in the art, and include considerations such as toxicity and forms that prevent the composition or formulation from exerting its effect.

By "systemic administration" is meant *in vivo* systemic absorption or accumulation of drugs in the blood stream followed by distribution throughout the entire body. Administration routes that lead to systemic absorption include, without limitation: intravenous, subcutaneous, intraperitoneal, inhalation, oral, intrapulmonary and intramuscular. Each of these administration routes exposes the siNA molecules of the invention to an accessible diseased tissue. The rate of entry of a drug into the circulation has been shown to be a function of molecular weight or size. The use of a liposome or other drug carrier comprising the compounds of the instant invention can potentially localize the drug, for example, in certain tissue types, such as the tissues of the reticular endothelial system (RES). A liposome formulation that can facilitate the association of drug with the surface of cells, such as, lymphocytes and macrophages is also useful. This approach can provide enhanced delivery of the drug to target cells by taking advantage of the specificity of macrophage and lymphocyte immune recognition of abnormal cells, such as cells producing excess MDR.

By "pharmaceutically acceptable formulation" is meant, a composition or formulation that allows for the effective distribution of the nucleic acid molecules of the instant invention in the physical location most suitable for their desired activity. Nonlimiting examples of agents suitable for formulation with the nucleic acid molecules of the instant invention include: P-glycoprotein inhibitors (such as Pluronic P85), which can enhance entry of drugs into the CNS (Jolliet-Riant and Tillement, 1999, Fundam. Clin. Pharmacol., 13, 16-26); biodegradable polymers, such as poly (DL-lactide-coglycolide) microspheres for sustained release delivery after intracerebral implantation (Emerich, DF et al, 1999, Cell Transplant, 8, 47-58) (Alkermes, Inc. Cambridge, MA); and loaded nanoparticles, such as those made of polybutylcyanoacrylate, which can deliver drugs across the blood brain barrier and can alter neuronal uptake mechanisms (Prog Neuropsychopharmacol Biol Psychiatry, 23, 941-949, 1999). Other non-limiting examples of delivery strategies for the nucleic acid molecules of the instant invention include material described in Boado et al., 1998, J. Pharm. Sci., 87, 1308-1315; Tyler et al., 1999, FEBS Lett., 421, 280-284; Pardridge et al., 1995, PNAS USA., 92, 5592-5596; Boado, 1995, Adv. Drug Delivery Rev., 15, 73-107; Aldrian-Herrada et al., 1998, Nucleic Acids Res., 26, 4910-4916; and Tyler et al., 1999, PNAS USA., 96, 7053-7058.

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The invention also features the use of the composition comprising surface-modified liposomes containing poly (ethylene glycol) lipids (PEG-modified, or long-circulating liposomes or stealth liposomes). These formulations offer a method for increasing the accumulation of drugs in target tissues. This class of drug carriers resists opsonization and elimination by the mononuclear phagocytic system (MPS or RES), thereby enabling longer blood circulation times and enhanced tissue exposure for the encapsulated drug (Lasic et al. Chem. Rev. 1995, 95, 2601-2627; Ishiwata et al., Chem. Pharm. Bull. 1995, 43, 1005-1011). Such liposomes have been shown to accumulate selectively in tumors, presumably by extravasation and capture in the neovascularized target tissues (Lasic et al., Science 1995, 267, 1275-1276; Oku et al., 1995, Biochim. Biophys. Acta, 1238, 86-90). The long-circulating liposomes enhance the pharmacokinetics and pharmacodynamics of DNA and RNA, particularly compared to conventional cationic liposomes which are known to accumulate in tissues of the MPS (Liu et al., J. Biol. Chem. 1995, 42, 24864-24870; Choi et al., International PCT Publication No. WO 96/10391; Ansell et al., International PCT Publication No. WO 96/10390; Holland et al.,

International PCT Publication No. WO 96/10392). Long-circulating liposomes are also likely to protect drugs from nuclease degradation to a greater extent compared to cationic liposomes, based on their ability to avoid accumulation in metabolically aggressive MPS tissues such as the liver and spleen.

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The present invention also includes compositions prepared for storage or administration that include a pharmaceutically effective amount of the desired compounds in a pharmaceutically acceptable carrier or diluent. Acceptable carriers or diluents for therapeutic use are well known in the pharmaceutical art, and are described, for example, in *Remington's Pharmaceutical Sciences*, Mack Publishing Co. (A.R. Gennaro edit. 1985), hereby incorporated by reference herein. For example, preservatives, stabilizers, dyes and flavoring agents can be provided. These include sodium benzoate, sorbic acid and esters of *p*-hydroxybenzoic acid. In addition, antioxidants and suspending agents can be used.

A pharmaceutically effective dose is that dose required to prevent, inhibit the occurrence, or treat (alleviate a symptom to some extent, preferably all of the symptoms) of a disease state. The pharmaceutically effective dose depends on the type of disease, the composition used, the route of administration, the type of mammal being treated, the physical characteristics of the specific mammal under consideration, concurrent medication, and other factors that those skilled in the medical arts will recognize. Generally, an amount between 0.1 mg/kg and 100 mg/kg body weight/day of active ingredients is administered dependent upon potency of the negatively charged polymer.

The nucleic acid molecules of the invention and formulations thereof can be administered orally, topically, parenterally, by inhalation or spray, or rectally in dosage unit formulations containing conventional non-toxic pharmaceutically acceptable carriers, adjuvants and/or vehicles. The term parenteral as used herein includes percutaneous, subcutaneous, intravascular (e.g., intravenous), intramuscular, or intrathecal injection or infusion techniques and the like. In addition, there is provided a pharmaceutical formulation comprising a nucleic acid molecule of the invention and a pharmaceutically acceptable carrier. One or more nucleic acid molecules of the invention can be present in association with one or more non-toxic pharmaceutically acceptable carriers and/or diluents and/or adjuvants, and if desired other active ingredients. The pharmaceutical

compositions containing nucleic acid molecules of the invention can be in a form suitable for oral use, for example, as tablets, troches, lozenges, aqueous or oily suspensions, dispersible powders or granules, emulsion, hard or soft capsules, or syrups or elixirs.

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Compositions intended for oral use can be prepared according to any method known to the art for the manufacture of pharmaceutical compositions and such compositions can contain one or more such sweetening agents, flavoring agents, coloring agents or preservative agents in order to provide pharmaceutically elegant and palatable preparations. Tablets contain the active ingredient in admixture with non-toxic pharmaceutically acceptable excipients that are suitable for the manufacture of tablets. These excipients can be, for example, inert diluents; such as calcium carbonate, sodium carbonate, lactose, calcium phosphate or sodium phosphate; granulating and disintegrating agents, for example, corn starch, or alginic acid; binding agents, for example starch, gelatin or acacia; and lubricating agents, for example magnesium stearate, stearic acid or talc. The tablets can be uncoated or they can be coated by known techniques. In some cases such coatings can be prepared by known techniques to delay disintegration and absorption in the gastrointestinal tract and thereby provide a sustained action over a longer period. For example, a time delay material such as glyceryl monosterate or glyceryl distearate can be employed.

Formulations for oral use can also be presented as hard gelatin capsules wherein the active ingredient is mixed with an inert solid diluent, for example, calcium carbonate, calcium phosphate or kaolin, or as soft gelatin capsules wherein the active ingredient is mixed with water or an oil medium, for example peanut oil, liquid paraffin or olive oil.

Aqueous suspensions contain the active materials in a mixture with excipients suitable for the manufacture of aqueous suspensions. Such excipients are suspending agents, for example sodium carboxymethylcellulose, methylcellulose, hydropropylmethylcellulose, sodium alginate, polyvinylpyrrolidone, gum tragacanth and gum acacia; dispersing or wetting agents can be a naturally-occurring phosphatide, for example, lecithin, or condensation products of an alkylene oxide with fatty acids, for example polyoxyethylene stearate, or condensation products of ethylene oxide with long chain aliphatic alcohols, for example heptadecaethyleneoxycetanol, or condensation products of ethylene oxide with partial esters derived from fatty acids and a hexitol such as

polyoxyethylene sorbitol monooleate, or condensation products of ethylene oxide with partial esters derived from fatty acids and hexitol anhydrides, for example polyethylene sorbitan monooleate. The aqueous suspensions can also contain one or more preservatives, for example ethyl, or n-propyl p-hydroxybenzoate, one or more coloring agents, one or more flavoring agents, and one or more sweetening agents, such as sucrose or saccharin.

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Oily suspensions can be formulated by suspending the active ingredients in a vegetable oil, for example arachis oil, olive oil, sesame oil or coconut oil, or in a mineral oil such as liquid paraffin. The oily suspensions can contain a thickening agent, for example beeswax, hard paraffin or cetyl alcohol. Sweetening agents and flavoring agents can be added to provide palatable oral preparations. These compositions can be preserved by the addition of an anti-oxidant such as ascorbic acid

Dispersible powders and granules suitable for preparation of an aqueous suspension by the addition of water provide the active ingredient in admixture with a dispersing or wetting agent, suspending agent and one or more preservatives. Suitable dispersing or wetting agents or suspending agents are exemplified by those already mentioned above. Additional excipients, for example sweetening, flavoring and coloring agents, can also be present.

Pharmaceutical compositions of the invention can also be in the form of oil-in-water emulsions. The oily phase can be a vegetable oil or a mineral oil or mixtures of these. Suitable emulsifying agents can be naturally-occurring gums, for example gum acacia or gum tragacanth, naturally-occurring phosphatides, for example soy bean, lecithin, and esters or partial esters derived from fatty acids and hexitol, anhydrides, for example sorbitan monooleate, and condensation products of the said partial esters with ethylene oxide, for example polyoxyethylene sorbitan monooleate. The emulsions can also contain sweetening and flavoring agents.

Syrups and elixirs can be formulated with sweetening agents, for example glycerol, propylene glycol, sorbitol, glucose or sucrose. Such formulations can also contain a demulcent, a preservative and flavoring and coloring agents. The pharmaceutical compositions can be in the form of a sterile injectable aqueous or oleaginous suspension. This suspension can be formulated according to the known art using those suitable

dispersing or wetting agents and suspending agents that have been mentioned above. The sterile injectable preparation can also be a sterile injectable solution or suspension in a non-toxic parentally acceptable diluent or solvent, for example as a solution in 1,3-butanediol. Among the acceptable vehicles and solvents that can be employed are water, Ringer's solution and isotonic sodium chloride solution. In addition, sterile, fixed oils are conventionally employed as a solvent or suspending medium. For this purpose, any bland fixed oil can be employed including synthetic mono-or diglycerides. In addition, fatty acids such as oleic acid find use in the preparation of injectables.

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The nucleic acid molecules of the invention can also be administered in the form of suppositories, e.g., for rectal administration of the drug. These compositions can be prepared by mixing the drug with a suitable non-irritating excipient that is solid at ordinary temperatures but liquid at the rectal temperature and will therefore melt in the rectum to release the drug. Such materials include cocoa butter and polyethylene glycols.

Nucleic acid molecules of the invention can be administered parenterally in a sterile medium. The drug, depending on the vehicle and concentration used, can either be suspended or dissolved in the vehicle. Advantageously, adjuvants such as local anesthetics, preservatives and buffering agents can be dissolved in the vehicle.

Dosage levels of the order of from about 0.1 mg to about 140 mg per kilogram of body weight per day are useful in the treatment of the above-indicated conditions (about 0.5 mg to about 7 g per subject per day). The amount of active ingredient that can be combined with the carrier materials to produce a single dosage form varies depending upon the host treated and the particular mode of administration. Dosage unit forms generally contain between from about 1 mg to about 500 mg of an active ingredient.

It is understood that the specific dose level for any particular subject depends upon a variety of factors including the activity of the specific compound employed, the age, body weight, general health, sex, diet, time of administration, route of administration, and rate of excretion, drug combination and the severity of the particular disease undergoing therapy.

For administration to non-human animals, the composition can also be added to the animal feed or drinking water. It can be convenient to formulate the animal feed and

drinking water compositions so that the animal takes in a therapeutically appropriate quantity of the composition along with its diet. It can also be convenient to present the composition as a premix for addition to the feed or drinking water.

The nucleic acid molecules of the present invention can also be administered to a subject in combination with other therapeutic compounds to increase the overall therapeutic effect. The use of multiple compounds to treat an indication can increase the beneficial effects while reducing the presence of side effects.

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In one embodiment, the invention comprises compositions suitable for administering nucleic acid molecules of the invention to specific cell types. For example, the asialoglycoprotein receptor (ASGPr) (Wu and Wu, 1987, J. Biol. Chem. 262, 4429-4432) is unique to hepatocytes and binds branched galactose-terminal glycoproteins, such as asialoorosomucoid (ASOR). In another example, the folate receptor is overexpressed in many cancer cells. Binding of such glycoproteins, synthetic glycoconjugates, or folates to the receptor takes place with an affinity that strongly depends on the degree of branching of the oligosaccharide chain, for example, triatennary structures are bound with greater affinity than biatenarry or monoatennary chains (Baenziger and Fiete, 1980, Cell, 22, 611-620; Connolly et al., 1982, J. Biol. Chem., 257, 939-945). Lee and Lee, 1987, Glycoconjugate J., 4, 317-328, obtained this high specificity through the use of N-acetyl-D-galactosamine as the carbohydrate moiety, which has higher affinity for the receptor, compared to galactose. This "clustering effect" has also been described for the binding and uptake of mannosyl-terminating glycoproteins or glycoconjugates (Ponpipom et al., 1981, J. Med. Chem., 24, 1388-1395). The use of galactose, galactosamine, or folate based conjugates to transport exogenous compounds across cell membranes can provide a targeted delivery approach to, for example, the treatment of liver disease, cancers of the liver, or other cancers. The use of bioconjugates can also provide a reduction in the required dose of therapeutic compounds required for treatment. Furthermore, therapeutic bioavialability, pharmacodynamics, and pharmacokinetic parameters can be modulated through the use of nucleic acid bioconjugates of the invention. Non-limiting examples of such bioconjugates are described in Vargeese et al., USSN 10/201,394, filed August 13, 2001; and Matulic-Adamic et al., USSN 60/362,016, filed March 6, 2002.

Alternatively, certain siNA molecules of the instant invention can be expressed within cells from eukaryotic promoters (e.g., Izant and Weintraub, 1985, Science, 229, 345; McGarry and Lindquist, 1986, Proc. Natl. Acad. Sci., USA 83, 399; Scanlon et al., 1991, Proc. Natl. Acad. Sci. USA, 88, 10591-5; Kashani-Sabet et al., 1992, Antisense Res. Dev., 2, 3-15; Dropulic et al., 1992, J. Virol., 66, 1432-41; Weerasinghe et al., 1991, J. Virol., 65, 5531-4; Ojwang et al., 1992, Proc. Natl. Acad. Sci. USA, 89, 10802-6; Chen et al., 1992, Nucleic Acids Res., 20, 4581-9; Sarver et al., 1990 Science, 247, 1222-1225; Thompson et al., 1995, Nucleic Acids Res., 23, 2259; Good et al., 1997, Gene Therapy, 4, 45. Those skilled in the art realize that any nucleic acid can be expressed in eukaryotic cells from the appropriate DNA/RNA vector. The activity of such nucleic acids can be augmented by their release from the primary transcript by a enzymatic nucleic acid (Draper et al., PCT WO 93/23569, and Sullivan et al., PCT WO 94/02595; Ohkawa et al., 1992, Nucleic Acids Symp. Ser., 27, 15-6; Taira et al., 1991, Nucleic Acids Res., 19, 5125-30; Ventura et al., 1993, Nucleic Acids Res., 21, 3249-55; Chowrira et al., 1994, J. Biol. Chem., 269, 25856.

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In another aspect of the invention, RNA molecules of the present invention can be expressed from transcription units (see for example Couture et al., 1996, TIG., 12, 510) inserted into DNA or RNA vectors. The recombinant vectors can be DNA plasmids or viral vectors siNA expressing viral vectors can be constructed based on, but not limited to, adeno-associated virus, retrovirus, adenovirus, or alphavirus. In another embodiment, pol III based constructs are used to express nucleic acid molecules of the invention (see for example Thompson, U.S. Pats. Nos. 5,902,880 and 6,146,886). The recombinant vectors capable of expressing the siNA molecules can be delivered as described above, and persist in target cells. Alternatively, viral vectors can be used that provide for transient expression of nucleic acid molecules. Such vectors can be repeatedly administered as necessary. Once expressed, the siNA molecule interacts with the target mRNA and generates an RNAi response. Delivery of siNA molecule expressing vectors can be systemic, such as by intravenous or intra-muscular administration, by administration to target cells ex-planted from a subject followed by reintroduction into the subject, or by any other means that would allow for introduction into the desired target cell (for a review see Couture et al., 1996, TIG., 12, 510).

In one aspect the invention features an expression vector comprising a nucleic acid sequence encoding at least one siNA molecule of the instant invention. The expression vector can encode one or both strands of a siNA duplex, or a single self-complementary strand that self hybridizes into a siNA duplex. The nucleic acid sequences encoding the siNA molecules of the instant invention can be operably linked in a manner that allows expression of the siNA molecule (see for example Paul et al., 2002, Nature Biotechnology, 19, 505; Miyagishi and Taira, 2002, Nature Biotechnology, 19, 497; Lee et al., 2002, Nature Biotechnology, 19, 500; and Novina et al., 2002, Nature Medicine, advance online publication doi:10.1038/nm725).

In another aspect, the invention features an expression vector comprising: a) a transcription initiation region (e.g., eukaryotic pol I, II or III initiation region); b) a transcription termination region (e.g., eukaryotic pol I, II or III termination region); and c) a nucleic acid sequence encoding at least one of the siNA molecules of the instant invention; wherein said sequence is operably linked to said initiation region and said termination region, in a manner that allows expression and/or delivery of the siNA molecule. The vector can optionally include an open reading frame (ORF) for a protein operably linked on the 5' side or the 3'-side of the sequence encoding the siNA of the invention; and/or an intron (intervening sequences).

Transcription of the siNA molecule sequences can be driven from a promoter for eukaryotic RNA polymerase I (pol I), RNA polymerase II (pol II), or RNA polymerase III (pol III). Transcripts from pol II or pol III promoters are expressed at high levels in all cells; the levels of a given pol II promoter in a given cell type depends on the nature of the gene regulatory sequences (enhancers, silencers, etc.) present nearby. Prokaryotic RNA polymerase promoters are also used, providing that the prokaryotic RNA polymerase enzyme is expressed in the appropriate cells (Elroy-Stein and Moss, 1990, *Proc. Natl. Acad. Sci. U S A*, 87, 6743-7; Gao and Huang 1993, *Nucleic Acids Res.*, 21, 2867-72; Lieber et al., 1993, *Methods Enzymol.*, 217, 47-66; Zhou et al., 1990, *Mol. Cell. Biol.*, 10, 4529-37). Several investigators have demonstrated that nucleic acid molecules expressed from such promoters can function in mammalian cells (e.g. Kashani-Sabet et al., 1992, *Antisense Res. Dev.*, 2, 3-15; Ojwang et al., 1992, *Proc. Natl. Acad. Sci. U S A*, 89, 10802-6; Chen et al., 1992, *Nucleic Acids Res.*, 20, 4581-9; Yu et al., 1993, *Proc. Natl. Acad. Sci. U S A*, 90, 6340-4; L'Huillier et al., 1992, *EMBO J.*, 11,

4411-8; Lisziewicz et al., 1993, Proc. Natl. Acad. Sci. U. S. A, 90, 8000-4; Thompson et al., 1995, Nucleic Acids Res., 23, 2259; Sullenger & Cech, 1993, Science, 262, 1566). More specifically, transcription units such as the ones derived from genes encoding U6 small nuclear (snRNA), transfer RNA (tRNA) and adenovirus VA RNA are useful in generating high concentrations of desired RNA molecules such as siNA in cells (Thompson et al., supra; Couture and Stinchcomb, 1996, supra; Noonberg et al., 1994, Nucleic Acid Res., 22, 2830; Noonberg et al., U.S. Pat. No. 5,624,803; Good et al., 1997, Gene Ther., 4, 45; Beigelman et al., International PCT Publication No. WO 96/18736. The above siNA transcription units can be incorporated into a variety of vectors for introduction into mammalian cells, including but not restricted to, plasmid DNA vectors, viral DNA vectors (such as adenovirus or adeno-associated virus vectors), or viral RNA vectors (such as retroviral or alphavirus vectors) (for a review see Couture and Stinchcomb, 1996, supra).

In another aspect the invention features an expression vector comprising a nucleic acid sequence encoding at least one of the siNA molecules of the invention in a manner that allows expression of that siNA molecule. The expression vector comprises in one embodiment; a) a transcription initiation region; b) a transcription termination region; and c) a nucleic acid sequence encoding at least one strand of the siNA molecule, wherein the sequence is operably linked to the initiation region and the termination region in a manner that allows expression and/or delivery of the siNA molecule.

In another embodiment the expression vector comprises: a) a transcription initiation region; b) a transcription termination region; c) an open reading frame; and d) a nucleic acid sequence encoding at least one strand of a siNA molecule, wherein the sequence is operably linked to the 3'-end of the open reading frame and wherein the sequence is operably linked to the initiation region, the open reading frame and the termination region in a manner that allows expression and/or delivery of the siNA molecule. In yet another embodiment, the expression vector comprises: a) a transcription initiation region; b) a transcription termination region; c) an intron; and d) a nucleic acid sequence encoding at least one siNA molecule, wherein the sequence is operably linked to the initiation region, the intron and the termination region in a manner which allows expression and/or delivery of the nucleic acid molecule.

In another embodiment, the expression vector comprises: a) a transcription initiation region; b) a transcription termination region; c) an intron; d) an open reading frame; and e) a nucleic acid sequence encoding at least one strand of a siNA molecule, wherein the sequence is operably linked to the 3'-end of the open reading frame and wherein the sequence is operably linked to the initiation region, the intron, the open reading frame and the termination region in a manner which allows expression and/or delivery of the siNA molecule.

Examples:

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The following are non-limiting examples showing the selection, isolation, synthesis and activity of nucleic acids of the instant invention.

Example 1: Tandem synthesis of siNA constructs

Exemplary siNA molecules of the invention are synthesized in tandem using a cleavable linker, for example, a succinyl-based linker. Tandem synthesis as described herein is followed by a one-step purification process that provides RNAi molecules in high yield. This approach is highly amenable to siNA synthesis in support of high throughput RNAi screening, and can be readily adapted to multi-column or multi-well synthesis platforms.

After completing a tandem synthesis of a siNA oligo and its complement in which the 5'-terminal dimethoxytrityl (5'-O-DMT) group remains intact (trityl on synthesis), the oligonucleotides are deprotected as described above. Following deprotection, the siNA sequence strands are allowed to spontaneously hybridize. This hybridization yields a duplex in which one strand has retained the 5'-O-DMT group while the complementary strand comprises a terminal 5'-hydroxyl. The newly formed duplex behaves as a single molecule during routine solid-phase extraction purification (Trityl-On purification) even though only one molecule has a dimethoxytrityl group. Because the strands form a stable duplex, this dimethoxytrityl group (or an equivalent group, such as other trityl groups or other hydrophobic moieties) is all that is required to purify the pair of oligos, for example, by using a C18 cartridge.

Standard phosphoramidite synthesis chemistry is used up to the point of introducing a tandem linker, such as an inverted deoxy abasic succinate or glyceryl succinate linker (see Figure 1) or an equivalent cleavable linker. A non-limiting example of linker coupling conditions that can be used includes a hindered base such as diisopropylethylamine (DIPA) and/or DMAP in the presence of an activator reagent such as Bromotripyrrolidinophosphoniumhexaflurorophosphate (PyBrOP). After the linker is coupled, standard synthesis chemistry is utilized to complete synthesis of the second sequence leaving the terminal the 5'-O-DMT intact. Following synthesis, the resulting oligonucleotide is deprotected according to the procedures described herein and quenched with a suitable buffer, for example with 50mM NaOAc or 1.5M NH₄H₂CO₃.

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Purification of the siNA duplex can be readily accomplished using solid phase extraction, for example using a Waters C18 SepPak 1g cartridge conditioned with 1 column volume (CV) of acetonitrile, 2 CV H2O, and 2 CV 50mM NaOAc. The sample is loaded and then washed with 1 CV H2O or 50mM NaOAc. Failure sequences are eluted with 1 CV 14% ACN (Aqueous with 50mM NaOAc and 50mM NaCl). The column is then washed, for example with 1 CV H2O followed by on-column detritylation, for example by passing 1 CV of 1% aqueous trifluoroacetic acid (TFA) over the column, then adding a second CV of 1% aqueous TFA to the column and allowing to stand for approximately 10 minutes. The remaining TFA solution is removed and the column washed with H2O followed by 1 CV 1M NaCl and additional H2O. The siNA duplex product is then eluted, for example, using 1 CV 20% aqueous CAN.

Figure 2 provides an example of MALDI-TOV mass spectrometry analysis of a purified siNA construct in which each peak corresponds to the calculated mass of an individual siNA strand of the siNA duplex. The same purified siNA provides three peaks when analyzed by capillary gel electrophoresis (CGE), one peak presumably corresponding to the duplex siNA, and two peaks presumably corresponding to the separate siNA sequence strands. Ion exchange HPLC analysis of the same siNA contract only shows a single peak. Testing of the purified siNA construct using a luciferase reporter assay described below demonstrated the same RNAi activity compared to siNA constructs generated from separately synthesized oligonucleotide sequence strands.

Example 2: Serum stability of chemically modified siNA constructs

Chemical modifications were introduced into siNA constructs to determine the stability of these constructs compared to native siNA oligonucleotides (containing two thymidine nucleotide overhangs) in human serum. An investigation of the serum stability of RNA duplexes revealed that siNA constructs consisting of all RNA nucleotides containing two thymidine nucleotide overhangs have a half-life in serum of 15 seconds, whereas chemically modified siNA constructs remained stable in serum for 1 to 3 days depending on the extent of modification. RNAi stability tests were performed by internally labeling one strand (strand 1) of siNA and duplexing with 1.5 X the concentration of the complementary siNA strand (strand 2) (to insure all labeled material was in duplex form). Duplexed siNA constructs were then tested for stability by incubating at a final concentration of 2µM siNA (strand 2 concentration) in 90% mouse or human serum for time-points of 30sec, 1min, 5min, 30min, 90min, 4hrs 10min, 16hrs 24min, and 49hrs. Time points were run on a 15% denaturing polyacrylamide gels and analyzed on a phosphoimager.

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Internal labeling was performed via kinase reactions with polynucleotide kinase (PNK) and ³²P-γ-ATP, with addition of radiolabeled phosphate at nucleotide 13 of strand 2, counting in from the 3' side. Ligation of the remaining 8-mer fragments with T4 RNA ligase resulted in the full length, 21-mer, strand 2. Duplexing of RNAi was done by adding appropriate concentrations of the siNA oligonucleotides and heating to 95° C for 5min followed by slow cooling to room temperature. Reactions were performed by adding 100% serum to the siNA duplexes and incubating at 37° C, then removing aliquots at desired time-points. Results of this study are summarized in Figure 3. As shown in the Figure 3, chemically modified siNA molecules (e.g., SEQ ID NOs: 925/927, 925/928, 925/929, 925/930, and 925/931) have significantly increased serum stability compared to an siNA construct having all ribonucleotides except a 3'-terminal dithymidine (TT) modification (e.g., SEQ ID NOs: 925/926).

Example 3: Identification of potential siNA target sites in any RNA sequence

The sequence of an RNA target of interest, such as a viral or human mRNA transcript, is screened for target sites, for example by using a computer folding algorithm. In a non-limiting example, the sequence of a gene or RNA gene transcript derived from a database, such as Genbank, is used to generate siNA targets having complementarity to

the target. Such sequences can be obtained from a database, or can be determined experimentally as known in the art. Target sites that are known, for example, those target sites determined to be effective target sites based on studies with other nucleic acid molecules, for example ribozymes or antisense, or those targets known to be associated with a disease or condition such as those sites containing mutations or deletions, can be used to design siNA molecules targeting those sites. Various parameters can be used to determine which sites are the most suitable target sites within the target RNA sequence. These parameters include but are not limited to secondary or tertiary RNA structure, the nucleotide base composition of the target sequence, the degree of homology between various regions of the target sequence, or the relative position of the target sequence within the RNA transcript. Based on these determinations, any number of target sites within the RNA transcript can be chosen to screen siNA molecules for efficacy, for example by using in vitro RNA cleavage assays, cell culture, or animal models. In a nonlimiting example, anywhere from 1 to 1000 target sites are chosen within the transcript based on the size of the siNA construct to be used. High throughput screening assays can be developed for screening siNA molecules using methods known in the art, such as with multi-well or multi-plate assays or combinatorial/siNA library screening assays to determine efficient reduction in target gene expression.

Example 4: Selection of siNA molecule target sites in a RNA

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The following non-limiting steps can be used to carry out the selection of siNAs targeting a given gene sequence or transcript.

The target sequence is parsed in silico into a list of all fragments or subsequences of a particular length, for example 23 nucleotide fragments, contained within the target sequence. This step is typically carried out using a custom Perl script, but commercial sequence analysis programs such as Oligo, MacVector, or the GCG Wisconsin Package can be employed as well.

In some instances the siNAs correspond to more than one target sequence; such would be the case for example in targeting different transcripts of the same gene, targeting different transcripts of more than one gene, or for targeting both the human gene and an animal homolog. In this case, a subsequence list of a particular length is generated for each of the targets, and then the lists are compared to find matching sequences in each

list. The subsequences are then ranked according to the number of target sequences that contain the given subsequence; the goal is to find subsequences that are present in most or all of the target sequences. Alternately, the ranking can identify subsequences that are unique to a target sequence, such as a mutant target sequence. Such an approach would enable the use of siNA to target specifically the mutant sequence and not effect the expression of the normal sequence.

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In some instances the siNA subsequences are absent in one or more sequences while present in the desired target sequence; such would be the case if the siNA targets a gene with a paralogous family member that is to remain untargeted. As in case 2 above, a subsequence list of a particular length is generated for each of the targets, and then the lists are compared to find sequences that are present in the target gene but are absent in the untargeted paralog.

The ranked siNA subsequences can be further analyzed and ranked according to GC content. A preference can be given to sites containing 30-70% GC, with a further preference to sites containing 40-60% GC.

The ranked siNA subsequences can be further analyzed and ranked according to self-folding and internal hairpins. Weaker internal folds are preferred; strong hairpin structures are to be avoided.

The ranked siNA subsequences can be further analyzed and ranked according to whether they have runs of GGG or CCC in the sequence. GGG (or even more Gs) in either strand can make oligonucleotide synthesis problematic and can potentially interfere with RNAi activity, so it is avoided whenever other appropriately suitable sequences are available. CCC is searched in the target strand because that will place GGG in the antisense strand.

The ranked siNA subsequences can be further analyzed and ranked according to whether they have the dinucleotide UU (uridine dinucleotide) on the 3'-end of the sequence, and/or AA on the 5'-end of the sequence (to yield 3' UU on the antisense sequence). These sequences allow one to design siNA molecules with terminal TT thymidine dinucleotides.

Four or five target sites are chosen from the ranked list of subsequences as described above. For example, in subsequences having 23 nucleotides, the right 21 nucleotides of each chosen 23-mer subsequence are then designed and synthesized for the upper (sense) strand of the siNA duplex, while the reverse complement of the left 21 nucleotides of each chosen 23-mer subsequence are then designed and synthesized for the lower (antisense) strand of the siNA duplex (see Tables I). If terminal TT residues are desired for the sequence (as described in paragraph 7), then the two 3' terminal nucleotides of both the sense and antisense strands are replaced by TT prior to synthesizing the oligos.

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The siNA molecules are screened in an in vitro, cell culture or animal model system to identify the most active siNA molecule or the most preferred target site within the target RNA sequence.

In an alternate approach, a pool of siNA constructs specific to a target sequence is used to screen for target sites in cells expressing target RNA, such as human HeLa cells. The general strategy used in this approach is shown in Figure 21. A non-limiting example of such as pool is a pool comprising sequences having antisense sequences complementary to the target RNA sequence and sense sequences complementary to the antisense sequences. Cells (e.g., HeLa cells) expressing the target gene are transfected with the pool of siNA constructs and cells that demonstrate a phenotype associated with gene silencing are sorted. The pool of siNA constructs can be chemically modified as described herein and synthesized, for example, in a high throughput manner. The siNA from cells demonstrating a positive phenotypic change (e.g., decreased target mRNA levels or target protein expression), are identified, for example by positional analysis within the assay, and are used to determine the most suitable target site(s) within the target RNA sequence based upon the complementary sequence to the corresponding siNA antisense strand identified in the assay.

Example 5: RNAi activity of chemically modified siNA constructs

Short interfering nucleic acid (siNA) is emerging as a powerful tool for gene regulation. All-ribose siNA duplexes activate the RNAi pathway but have limited utility as therapeutic compounds due to their nuclease sensitivity and short half-life in serum, as shown in Example 2 above. To develop nuclease-resistant siNA constructs for *in vivo*

applications, siNAs that target luciferase mRNA and contain stabilizing chemical modifications were tested for activity in HeLa cells. The sequences for the siNA oligonucleotide sequences used in this study are shown in **Table I**. Modifications included phosphorothioate linkages (P=S), 2'-O-methyl nucleotides, or 2'-fluoro (F) nucleotides in one or both siNA strands and various 3'-end stabilization chemistries, including 3'-glyceryl, 3'-inverted abasic, 3'-inverted Thymidine, and/or Thymidine. Active siNA containing stabilizing modifications such as described herein should prove useful for *in vivo* applications.

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A luciferase reporter system was utilized to test RNAi activity of chemically modified siNA constructs compared to siNA constructs consisting of all RNA nucleotides containing two thymidine nucleotide overhangs. Sense and antisense siNA strands (20 uM each) were annealed by incubation in buffer (100 mM potassium acetate, 30 mM HEPES-KOH, pH 7.4, 2 mM magnesium acetate) for 1 min. at 90°C followed by 1 hour at 37°C. Plasmids encoding firefly luciferase (pGL2) and renilla luciferase (pRLSV40) were purchased from Promega Biotech.

HeLa S3 cells were grown at 37°C in DMEM with 5% FBS and seeded at 15,300 cells in 100 ul media per well of a 96-well plate 24 hours prior to transfection. For transfection, 4 ul Lipofectamine 2000 (Life Technologies) was added to 96 ul OPTI-MEM, vortexed and incubated at room temperature for 5 minutes. The 100 ul diluted lipid was then added to a microtiter tube containing 5 ul pGL2 (200ng/ul), 5 ul pRLSV40 (8 ng/ul) 6 ul siNA (25 nM or 10 nM final), and 84 ul OPTI-MEM, vortexed briefly and incubated at room temperature for 20 minutes. The transfection mix was then mixed briefly and 50 ul was added to each of three wells that contained HeLa S3 cells in 100 ul media. Cells were incubated for 20 hours after transfection and analyzed for luciferase expression using the Dual luciferase assay according to the manufacturer's instructions (Promega Biotech). The results of this study are summarized in Figures 4-16. The sequences of the siNA strands used in this study are shown in Table I and are referred to by RPI# in the figures. Normalized luciferase activity is reported as the ratio of firefly luciferase activity to renilla luciferase activity in the same sample. Error bars represent standard deviation of triplicate transfections. As shown in Figures 4-16, the RNAi activity of chemically modified constructs is comparable to that of control siNA constructs, which consist of all ribonucleotides at every position except the 3'-terminus

which comprises two thymidine nucleotide overhangs. In some instances, the RNAi activity of the chemically modified constructs is greater than the siNA construct consisting of all ribonucleotides at every position except the 3'-terminus which comprises two thymidine nucleotide overhangs. For example, Figure 4 shows results obtained from a screen using phosphorothioate modified siNA constructs; the RPI 27654/27659 construct contains phosphorothioate substitutions for every pyrimidine nucleotide in both sequences, the RPI 27657/27662 construct contains 5 terminal 3'-phosphorothioate substitutions in each strand, the RPI 27649/27658 construct contains all phosphorothioate substitutions only in the antisense strand, whereas the RPI 27649/27660 and RPI 27649/27661 constructs have unmodified sense strands and varying degrees of phosphorothioate substitutions in the antisense strand. All of these constructs show significant RNAi activity when compared to a scrambled siNA.

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Figure 5 shows results obtained from a screen using phosphorothicate (RPI 28253/28255 and RPI 28254/28256) and universal base substitutions (RPI 28257/28259 and RPI 28258/28260) compared to the same controls described above. As shown, these modifications show equivalent or better RNAi activity when compared to the control siNA construct.

Figure 6 shows results obtained from a screen using 2'-O-methyl modified siNA constructs in which the sense strand contains either 10 (RPI 28244/27650) or 5 (RPI 28245/27650) 2'-O-methyl substitutions, both with comparable activity to the control siNA construct.

Figure 7 shows results obtained from a screen using 2'-O-methyl or 2'-deoxy-2'-fluoro modified siNA constructs compared to a control construct consisting of all ribonucleotides at every position except the 3'-terminus which comprises two thymidine nucleotide overhangs.

Figure 8 compares a siNA construct containing six phosphorothioate substitutions in each strand (RPI 28460/28461), where 5 phosphorothioates are present at the 3' end and a single phosphorothioate is present at the 5' end of each strand. This motif shows very similar activity to the control siNA construct consisting of all ribonucleotides at every position except the 3'-terminus which comprises two thymidine nucleotide overhangs.

Figure 9 compares a siNA construct synthesized by the method of the invention described in Example 1, wherein an inverted deoxyabasic succinate linker was used to generate a siNA having a 3'-inverted deoxyabasic cap on the antisense strand of the siNA. This construct shows improved activity compared to the control siNA (siGL2) construct consisting of all ribonucleotides at every position except the 3'-terminus which comprises two thymidine nucleotide overhangs.

Figure 10 shows the results of an RNAi activity screen of chemically modified siNA constructs including 3'-glyceryl modified siNA constructs compared to an all RNA control siNA construct using a luciferase reporter system. These chemically modified siNAs were compared in the luciferase assay described herein at 1 nM and 10nM concentration using an all RNA siNA control (siGL2) having having 3'-terminal dithymidine (TT) and its corresponding inverted control (Inv siGL2). The background level of luciferase expression in the HeLa cells is designated by the "cells" column. Sense and antisense strands of chemically modified siNA constructs are shown by RPI number (sense strand/antisense strand). Sequences correspoding to these RPI numbers are shown in Table I. As shown in the Figure, the 3'-terminal modified siNA constructs retain significant RNAi activity compared to the control siNA (siGL2) construct.

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Figure 11 shows the results of an RNAi activity screen of chemically modified siNA constructs. The screen compared various combinations of sense strand chemical modifications and antisense strand chemical modifications. These chemically modified siNAs were compared in the luciferase assay described herein at 1 nM and 10nM concentration using an all RNA siNA control (siGL2) having having 3'-terminal dithymidine (TT) and its corresponding inverted control (Inv siGL2). The background level of luciferase expression in the HeLa cells is designated by the "cells" column. Sense and antisense strands of chemically modified siNA constructs are shown by RPI number (sense strand/antisense strand). Sequences correspoding to these RPI numbers are shown in Table I. As shown in the figure, the chemically modified RPI 30063/30430, RPI 30433/30430, and RPI 30063/30224 constructs retain significant RNAi activity compared to the control siNA construct. It should be noted that RPI 30433/30430 is a siNA construct having no ribonucleotides which retains significant RNAi activity compared to the constrol siGL2 construct in vitro, therefore, this construct is expected to

have both similar RNAi activity and improved stability compared to siNA constructs having ribonucleotides in vivo.

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Figure 12 shows the results of an RNAi activity screen of chemically modifed siNA constructs. The screen compared various combinations of sense strand chemical modifications and antisense strand chemical modifications. These chemically modified siNAs were compared in the luciferase assay described herein at 1 nM and 10nM concentration using an all RNA siNA control (siGL2) having having 3'-terminal dithymidine (TT) and its corresponding inverted control (Inv siGL2). The background level of luciferase expression in the HeLa cells is designated by the "cells" column. Sense and antisense strands of chemically modified siNA constructs are shown by RPI number (sense strand/antisense strand). Sequences correspoding to these RPI numbers are shown in Table I. As shown in the figure, the chemically modified RPI 30063/30224 and RPI 30063/30430 constructs retain significant RNAi activity compared to the control siNA (siGL2) construct. In addition, the antisense strand alone (RPI 30430) and an inverted control (RPI 30227/30229, having matched chemistry to RPI 30063/30224) were compared to the siNA duplexes described above. The antisense strand (RPI 30430) alone provides far less inhibition compared to the siNA duplexes using this sequence.

Figure 13 shows the results of an RNAi activity screen of chemically modified siNA constructs. The screen compared various combinations of sense strand chemical modifications and antisense strand chemical modifications. These chemically modified siNAs were compared in the luciferase assay described herein at 1 nM and 10nM concentration using an all RNA siNA control (siGL2) having having 3'-terminal dithymidine (TT) and its corresponding inverted control (Inv siGL2). The background level of luciferase expression in the HeLa cells is designated by the "cells" column. Sense and antisense strands of chemically modified siNA constructs are shown by RPI number (sense strand/antisense strand). Sequences corresponding to these RPI numbers are shown in Table I. In addition, an inverted control (RPI 30226/30229, having matched chemistry to RPI 30222/30224) was compared to the siNA duplexes described above. As shown in the figure, the chemically modified RPI 28251/30430, RPI 28251/30224, and RPI 30222/30224 constructs retain significant RNAi activity compared to the control siNA construct, and the chemically modified RPI 28251/30430 construct demonstrates improved activity compared to the control siNA (siGL2) construct.

Figure 14 shows the results of an RNAi activity screen of chemically modified siNA constructs including various 3'-terminal modified siNA constructs compared to an all RNA control siNA construct using a luciferase reporter system. These chemically modified siNAs were compared in the luciferase assay described herein at 1 nM and 10nM concentration using an all RNA siNA control (siGL2) having having 3'-terminal dithymidine (TT) and its corresponding inverted control (Inv siGL2). The background level of luciferase expression in the HeLa cells is designated by the "cells" column. Sense and antisense strands of chemically modified siNA constructs are shown by RPI number (sense strand/antisense strand). Sequences correspoding to these RPI numbers are shown in Table I. As shown in the figure, the chemically modified RPI 30222/30546, 30222/30224, 30222/30551, 30222/30557 and 30222/30558 constructs retain significant RNAi activity compared to the control siNA construct.

Figure 15 shows the results of an RNAi activity screen of chemically modified siNA constructs. The screen compared various combinations of sense strand chemistries compared to a fixed antisense strand chemistry. These chemically modified siNAs were compared in the luciferase assay described herein at 1 nM and 10nM concentration using an all RNA siNA control (siGL2) having having 3'-terminal dithymidine (TT) and its corresponding inverted control (Inv siGL2). The background level of luciferase expression in the HeLa cells is designated by the "cells" column. Sense and antisense strands of chemically modified siNA constructs are shown by RPI number (sense strand/antisense strand). Sequences correspoding to these RPI numbers are shown in Table I. As shown in the figure, the chemically modified RPI 30063/30430, 30434/30430, and 30435/30430 constructs all demonstrate greater activity compared to the control siNA (siGL2) construct.

25 Example 6: RNAi activity titration

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A titration assay was performed to determine the lower range of siNA concentration required for RNAi activity both in a control siNA construct consisting of all RNA nucleotides containing two thymidine nucleotide overhangs and a chemically modified siNA construct comprising 5 phosphorothicate internucleotide linkages in both the sense and antisense strands. The assay was performed as described above, however, the siNA constructs were diluted to final concentrations between 2.5 nM and 0.025 nM. Results

are shown in **Figure 16**. As shown in **Figure 16**, the chemically modified siNA construct shows a very similar concentration dependent RNAi activity profile to the control siNA construct when compared to an inverted siNA sequence control.

Example 7: siNA design

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siNA target sites were chosen by analyzing sequences of the target RNA and optionally prioritizing the target sites on the basis of folding (structure of any given sequence analyzed to determine siNA accessibility to the target), by using a library of siNA molecules as described in Example 4, or alternately by using an *in vitro* siNA system as described in Example 9 herein. siNA molecules were designed that could bind each target and are optionally individually analyzed by computer folding to assess whether the siNA molecule can interact with the target sequence. Varying the length of the siNA molecules can be chosen to optimize activity. Generally, a sufficient number of complementary nucleotide bases are chosen to bind to, or otherwise interact with, the target RNA, but the degree of complementarity can be modulated to accommodate siNA duplexes or varying length or base composition. By using such methodologies, siNA molecules can be designed to target sites within any known RNA sequence, for example those RNA sequences corresponding to the any gene transcript.

Chemically modified siNA constructs are designed to provide nuclease stability for systemic administration in vivo and/or improved pharmacokinetic, localization, and delivery properties while preserving the ability to mediate RNAi activity. Chemical modifications as described herein are introduced synthetically using synthetic methods described herein and those generally known in the art. The synthetic siNA constructs are then assayed for nuclease stability in serum and/or cellular/tissue extracts (e.g. liver extracts). The synthetic siNA constructs are also tested in parallel for RNAi activity using an appropriate assay, such as a luciferase reporter assay as described herein or another suitable assay that can quantity RNAi activity. Synthetic siNA constructs that possess both nuclease stability and RNAi activity can be further modified and reevaluated in stability and activity assays. The chemical modifications of the stabilized active siNA constructs can then be applied to any siNA sequence targeting any chosen RNA and used, for example, in target screening assays to pick lead siNA compounds for therapeutic development (see for example Figure 24).

Example 8: Chemical Synthesis and Purification of siNA

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siNA molecules can be designed to interact with various sites in the RNA message, for example, target sequences within the RNA sequences described herein. The sequence of one strand of the siNA molecule(s) is complementary to the target site sequences described above. The siNA molecules can be chemically synthesized using methods described herein. Inactive siNA molecules that are used as control sequences can be synthesized by scrambling the sequence of the siNA molecules such that it is not complementary to the target sequence. Generally, siNA constructs can by synthesized using solid phase oligonucleotide synthesis methods as described herein (see for example Usman *et al.*, US Patent Nos. 5,804,683; 5,831,071; 5,998,203; 6,117,657; 6,353,098; 6,362,323; 6,437,117; 6,469,158; Scaringe *et al.*, US Patent Nos. 6,111,086; 6,008,400; 6,111,086 all incorporated by reference herein in their entirety).

In a non-limiting example, RNA oligonucleotides are synthesized in a stepwise fashion using the phosphoramidite chemistry as is known in the art. Standard phosphoramidite chemistry involves the use of nucleosides comprising any of 5'-O-dimethoxytrityl, 2'-O-tert-butyldimethylsilyl, 3'-O-2-Cyanoethyl N,N-diisopropylphosphoroamidite groups, and exocyclic amine protecting groups (e.g. N6-benzoyl adenosine, N4 acetyl cytidine, and N2-isobutyryl guanosine). Alternately, 2'-O-Silyl Ethers can be used in conjunction with acid-labile 2'-O-orthoester protecting groups in the synthesis of RNA as described by Scaringe supra. Differing 2' chemistries can require different protecting groups, for example 2'-deoxy-2'-amino nucleosides can utilize N-phthaloyl protection as described by Usman et al., US Patent 5,631,360, incorporated by reference herein in its entirety).

During solid phase synthesis, each nucleotide is added sequentially (3'- to 5'-direction) to the solid support-bound oligonucleotide. The first nucleoside at the 3'-end of the chain is covalently attached to a solid support (e.g., controlled pore glass or polystyrene) using various linkers. The nucleotide precursor, a ribonucleoside phosphoramidite, and activator are combined resulting in the coupling of the second nucleoside phosphoramidite onto the 5'-end of the first nucleoside. The support is then washed and any unreacted 5'-hydroxyl groups are capped with a capping reagent such as acetic anhydride to yield inactive 5'-acetyl moieties. The trivalent phosphorus linkage is

then oxidized to a more stable phosphate linkage. At the end of the nucleotide addition cycle, the 5'-O-protecting group is cleaved under suitable conditions (e.g., acidic conditions for trityl-based groups and Fluoride for silyl-based groups). The cycle is repeated for each subsequent nucleotide.

Modification of synthesis conditions can be used to optimize coupling efficiency, for example by using differing coupling times, differing reagent/phosphoramidite concentrations, differing contact times, differing solid supports and solid support linker chemistries depending on the particular chemical composition of the siNA to be synthesized. Deprotection and purification of the siNA can be performed as is generally described in Usman et al., US 5,831,071, US 6,353,098, US 6,437,117, and Bellon et al., US 6,054,576, US 6,162,909, US 6,303,773, incorporated by reference herein in their entirety or Scaringe *supra*,. Additionally, deprotection conditions can be modified to provide the best possible yield and purity of siNA constructs. For example, applicant has observed that oligonucleotides comprising 2'-deoxy-2'-fluoro nucleotides can degrade under inappropriate deprotection conditions. Such oligonucleotides are deprotected using aqueous methylamine at about 35°C for 30 minutes. If the 2'-deoxy-2'-fluoro containing oligonucleotide also comprises ribonucleotides, after deprotection with aqueous methylamine at about 35°C for 30 minutes, TEA-HF is added and the reaction maintained at about 65°C for an additional 15 minutes.

20 Example 9: RNAi in vitro assay to assess siNA activity

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An in vitro assay that recapitulates RNAi in a cell free system is used to evaluate siNA constructs specific to target RNA. The assay comprises the system described by Tuschl et al., 1999, Genes and Development, 13, 3191-3197 and Zamore et al., 2000, Cell, 101, 25-33 adapted for use with target RNA. A Drosophila extract derived from syncytial blastoderm is used to reconstitute RNAi activity in vitro. Target RNA is generated via in vitro transcription from an appropriate plasmid using T7 RNA polymerase or via chemical synthesis as described herein. Sense and antisense siNA strands (for example 20 uM each) are annealed by incubation in buffer (such as 100 mM potassium acetate, 30 mM HEPES-KOH, pH 7.4, 2 mM magnesium acetate) for 1 min. at 90°C followed by 1 hour at 37°C, then diluted in lysis buffer (for example 100 mM potassium acetate, 30 mM HEPES-KOH at pH 7.4, 2mM magnesium acetate). Annealing

can be monitored by gel electrophoresis on an agarose gel in TBE buffer and stained with ethidium bromide. The Drosophila lysate is prepared using zero to two-hour-old embryos from Oregon R flies collected on yeasted molasses agar that are dechorionated and lysed. The lysate is centrifuged and the supernatant isolated. The assay comprises a reaction mixture containing 50% lysate [vol/vol], RNA (10-50 pM final concentration), and 10% [vol/vol] lysis buffer containing siNA (10 nM final concentration). The reaction mixture also contains 10 mM creatine phosphate, 10 ug.ml creatine phosphokinase, 100 um GTP, 100 uM UTP, 100 uM CTP, 500 uM ATP, 5 mM DTT, 0.1 U/uL RNasin (Promega), and 100 uM of each amino acid. The final concentration of potassium acetate is adjusted to 100 mM. The reactions are pre-assembled on ice and preincubated at 25° C for 10 minutes before adding RNA, then incubated at 25° C for an additional 60 minutes. Reactions are quenched with 4 volumes of 1.25 x Passive Lysis Buffer (Promega). Target RNA cleavage is assayed by RT-PCR analysis or other methods known in the art and are compared to control reactions in which siNA is omitted from the reaction.

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Alternately, internally-labeled target RNA for the assay is prepared by *in vitro* transcription in the presence of [alpha-³²p] CTP, passed over a G 50 Sephadex column by spin chromatography and used as target RNA without further purification. Optionally, target RNA is 5'-³²P-end labeled using T4 polynucleotide kinase enzyme. Assays are performed as described above and target RNA and the specific RNA cleavage products generated by RNAi are visualized on an autoradiograph of a gel. The percentage of cleavage is determined by Phosphor Imager[®] quantitation of bands representing intact control RNA or RNA from control reactions without siNA and the cleavage products generated by the assay.

In one embodiment, this assay is used to determine target sites the RNA target for siNA mediated RNAi cleavage, wherein a plurality of siNA constructs are screened for RNAi mediated cleavage of the RNA target, for example, by analyzing the assay reaction by electrophoresis of labeled target RNA, or by northern blotting, as well as by other methodology well known in the art.

Example 10: Nucleic acid inhibition of target RNA in vivo

siNA molecules targeted to the target RNA are designed and synthesized as described above. These nucleic acid molecules can be tested for cleavage activity *in vivo*, for example, using the following procedure.

Two formats are used to test the efficacy of siNAs targeting a particular gene transcipt. First, the reagents are tested on target expressing cells (e.g., HeLa), to determine the extent of RNA and protein inhibition. siNA reagents are selected against the RNA target. RNA inhibition is measured after delivery of these reagents by a suitable transfection agent to cells. Relative amounts of target RNA are measured versus actin using real-time PCR monitoring of amplification (eg., ABI 7700 Taqman®). A comparison is made to a mixture of oligonucleotide sequences made to unrelated targets or to a randomized siNA control with the same overall length and chemistry, but randomly substituted at each position. Primary and secondary lead reagents are chosen for the target and optimization performed. After an optimal transfection agent concentration is chosen, a RNA time-course of inhibition is performed with the lead siNA molecule. In addition, a cell-plating format can be used to determine RNA inhibition.

Delivery of siNA to Cells

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Cells (e.g., HeLa) are seeded, for example, at 1x10⁵ cells per well of a six-well dish in EGM-2 (BioWhittaker) the day before transfection. siNA (final concentration, for example 20nM) and cationic lipid (e.g., final concentration 2µg/ml) are complexed in EGM basal media (Biowhittaker) at 37°C for 30 mins in polystyrene tubes. Following vortexing, the complexed siNA is added to each well and incubated for the times indicated. For initial optimization experiments, cells are seeded, for example, at 1x10³ in 96 well plates and siNA complex added as described. Efficiency of delivery of siNA to cells is determined using a fluorescent siNA complexed with lipid. Cells in 6-well dishes are incubated with siNA for 24 hours, rinsed with PBS and fixed in 2% paraformaldehyde for 15 minutes at room temperature. Uptake of siNA is visualized using a fluorescent microscope.

Taqman and Lightcycler quantification of mRNA

Total RNA is prepared from cells following siNA delivery, for example, using Qiagen RNA purification kits for 6-well or Rneasy extraction kits for 96-well assays. For

Taqman analysis, dual-labeled probes are synthesized with the reporter dye, FAM or JOE, covalently linked at the 5'-end and the quencher dye TAMRA conjugated to the 3'-end. One-step RT-PCR amplifications are performed on, for example, an ABI PRISM 7700 Sequence Detector using 50 µl reactions consisting of 10 µl total RNA, 100 nM forward primer, 900 nM reverse primer, 100 nM probe, 1X TaqMan PCR reaction buffer (PE-Applied Biosystems), 5.5 mM MgCl₂, 300 µM each dATP, dCTP, dGTP, and dTTP, 10U RNase Inhibitor (Promega), 1.25U AmpliTaq Gold (PE-Applied Biosystems) and 10U M-MLV Reverse Transcriptase (Promega). The thermal cycling conditions can consist of 30 min at 48°C, 10 min at 95°C, followed by 40 cycles of 15 sec at 95°C and 1 min at 60°C. Quantitation of mRNA levels is determined relative to standards generated from serially diluted total cellular RNA (300, 100, 33, 11 ng/rxn) and normalizing to \(\beta\)-actin or GAPDH mRNA in parallel TaqMan reactions. For each gene of interest an upper and lower primer and a fluorescently labeled probe are designed. Real time incorporation of SYBR Green I dye into a specific PCR product can be measured in glass capillary tubes using a lightcyler. A standard curve is generated for each primer pair using control cRNA. Values are represented as relative expression to GAPDH in each sample.

Western blotting

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Nuclear extracts can be prepared using a standard micro preparation technique (see for example Andrews and Faller, 1991, *Nucleic Acids Research*, 19, 2499). Protein extracts from supernatants are prepared, for example using TCA precipitation. An equal volume of 20% TCA is added to the cell supernatant, incubated on ice for 1 hour and pelleted by centrifugation for 5 minutes. Pellets are washed in acetone, dried and resuspended in water. Cellular protein extracts are run on a 10% Bis-Tris NuPage (nuclear extracts) or 4-12% Tris-Glycine (supernatant extracts) polyacrylamide gel and transferred onto nitro-cellulose membranes. Non-specific binding can be blocked by incubation, for example, with 5% non-fat milk for 1 hour followed by primary antibody for 16 hour at 4°C. Following washes, the secondary antibody is applied, for example (1:10,000 dilution) for 1 hour at room temperature and the signal detected with SuperSignal reagent (Pierce).

Example 11: Animal Models

Various animal models can be used to screen siNA constructs in vivo as are known in the art, for example those animal models that are used to evaluate other nucleic acid technologies such as enzymatic nucleic acid molecules (ribozymes) and/or antisense. Such animal models are used to test the efficacy of siNA molecules described herein. In a non-limiting example, siNA molecules that are designed as anti-angiogenic agents can be screened animal models. There are several animal models in which the antiangiogenesis effect of nucleic acids of the present invention, such as siNA, directed against genes associated with angiogenesis and/or metastais, such as VEGFR (e.g., VEGFR1, VEGFR2, and VEGFR3) genes. Typically a corneal model has been used to study angiogenesis in rat and rabbit since recruitment of vessels can easily be followed in this normally avascular tissue (Pandey et al., 1995 Science 268: 567-569). In these models, a small Teflon or Hydron disk pretreated with an angiogenesis factor (e.g. bFGF or VEGF) is inserted into a pocket surgically created in the comea. Angiogenesis is monitored 3 to 5 days later. siNA molecules directed against VEGFR mRNAs are delivered in the disk as well, or dropwise to the eye over the time course of the experiment. In another eye model, hypoxia has been shown to cause both increased expression of VEGF and neovascularization in the retina (Pierce et al., 1995 Proc. Natl. Acad. Sci. USA. 92: 905-909; Shweiki et al., 1992 J. Clin. Invest. 91: 2235-2243).

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Several animal models exist for screening of anti-angiogenic agents. These include corneal vessel formation following corneal injury (Burger et al., 1985 Cornea 4: 35-41; Lepri, et al., 1994 J. Ocular Pharmacol. 10: 273-280; Ormerod et al., 1990 Am. J. Pathol. 137: 1243-1252) or intracorneal growth factor implant (Grant et al., 1993 Diabetologia 36: 282-291; Pandey et al. 1995 supra; Zieche et al., 1992 Lab. Invest. 67: 711-715), vessel growth into Matrigel matrix containing growth factors (Passaniti et al., 1992 supra), female reproductive organ neovascularization following hormonal manipulation (Shweiki et al., 1993 Clin. Invest. 91: 2235-2243), several models involving inhibition of tumor growth in highly vascularized solid tumors (O'Reilly et al., 1994 Cell 79: 315-328; Senger et al., 1993 Cancer and Metas. Rev. 12: 303-324; Takahasi et al., 1994 Cancer Res. 54: 4233-4237; Kim et al., 1993 supra), and transient hypoxia-induced neovascularization in the mouse retina (Pierce et al., 1995 Proc. Natl. Acad. Sci. USA. 92: 905-909).gene

The cornea model, described in Pandey et al. supra, is the most common and well characterized anti-angiogenic agent efficacy screening model. This model involves an avascular tissue into which vessels are recruited by a stimulating agent (growth factor, thermal or alkalai burn, endotoxin). The corneal model would utilize the intrastromal corneal implantation of a Teflon pellet soaked in a VEGF-Hydron solution to recruit blood vessels toward the pellet which can be quantitated using standard microscopic and image analysis techniques. To evaluate their anti-angiogenic efficacy, ribozymes are applied topically to the eye or bound within Hydron on the Teflon pellet itself. This avascular cornea as well as the Matrigel model provide for low background assays. While the corneal model has been performed extensively in the rabbit, studies in the rat have also been conducted.

The mouse model (Passaniti et al., *supra*) is a non-tissue model which utilizes Matrigel, an extract of basement membrane (Kleinman et al., 1986) or Millipore[®] filter disk, which can be impregnated with growth factors and anti-angiogenic agents in a liquid form prior to injection. Upon subcutaneous administration at body temperature, the Matrigel or Millipore[®] filter disk forms a solid implant. VEGF embedded in the Matrigel or Millipore[®] filter disk is used to recruit vessels within the matrix of the Matrigel or Millipore[®] filter disk which can be processed histologically for endothelial cell specific vWF (factor VIII antigen) immunohistochemistry, Trichrome-Masson stain, or hemoglobin content. Like the cornea, the Matrigel or Millipore[®] filter disk are avascular; however, it is not tissue. In the Matrigel or Millipore[®] filter disk model, siNA molecules are administered within the matrix of the Matrigel or Millipore[®] filter disk to test their anti-angiogenic efficacy. Thus, delivery issues in this model, as with delivery of siNA molecules by Hydron- coated Teflon pellets in the rat cornea model, may be less problematic due to the homogeneous presence of the siNA within the respective matrix.

The Lewis lung carcinoma and B-16 murine melanoma models are well accepted models of primary and metastatic cancer and are used for initial screening of anti-cancer agents. These murine models are not dependent upon the use of immunodeficient mice, are relatively inexpensive, and minimize housing concerns. Both the Lewis lung and B-16 melanoma models involve subcutaneous implantation of approximately 106 tumor cells from metastatically aggressive tumor cell lines (Lewis lung lines 3LL or D122, LLc-

LN7; B-16-BL6 melanoma) in C57BL/6J mice. Alternatively, the Lewis lung model can be produced by the surgical implantation of tumor spheres (approximately 0.8 mm in diameter). Metastasis also may be modeled by injecting the tumor cells directly *i.v.*. In the Lewis lung model, microscopic metastases can be observed approximately 14 days following implantation with quantifiable macroscopic metastatic tumors developing within 21-25 days. The B-16 melanoma exhibits a similar time course with tumor neovascularization beginning 4 days following implantation. Since both primary and metastatic tumors exist in these models after 21-25 days in the same animal, multiple measurements can be taken as indices of efficacy. Primary tumor volume and growth latency as well as the number of micro- and macroscopic metastatic lung foci or number of animals exhibiting metastases can be quantitated. The percent increase in lifespan can also be measured. Thus, these models provide suitable primary efficacy assays for screening systemically administered siNA molecules and siNA formulations.

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In the Lewis lung and B-16 melanoma models, systemic pharmacotherapy with a wide variety of agents usually begins 1-7 days following tumor implantation/inoculation with either continuous or multiple administration regimens. Concurrent pharmacokinetic studies can be performed to determine whether sufficient tissue levels of siNA can be achieved for pharmacodynamic effect to be expected. Furthermore, primary tumors and secondary lung metastases can be removed and subjected to a variety of *in vitro* studies (*i.e.* target RNA reduction).

In utilizing these models to assess siNA activity, VEGFR1, VEGFR2, and/or VEGFR3 protein levels can be measured clinically or experimentally by FACS analysis. VEGFR1, VEGFR2, and/or VEGFR3 encoded mRNA levels will be assessed by Northern analysis, RNase-protection, primer extension analysis and/or quantitative RT-PCR. siNA molecules that block VEGFR1, VEGFR2, and/or VEGFR3 protein encoding mRNAs and therefore result in decreased levels of VEGFR1, VEGFR2, and/or VEGFR3 activity by more than 20% in vitro can be thus identified.

Example 12: siNA-mediated inhibition of angiogenesis in vivo

The purpose of this study was to assess the anti-angiogenic activity of siNA targeted against VEGFR1 in the rat comea model of VEGF induced angiogenesis (see above). These siNA molecules have matched inverted controls which are inactive since

they are not able to interact with the RNA target. The siNA molecules and VEGF were co-delivered using the filter disk method: Nitrocellulose filter disks (Millipore®) of 0.057 diameter were immersed in appropriate solutions and were surgically implanted in rat cornea as described by Pandey et al., supra.

The stimulus for angiogenesis in this study was the treatment of the filter disk with 30 µM VEGF which is implanted within the cornea's stroma. This dose yields reproducible neovascularization stemming from the pericorneal vascular plexus growing toward the disk in a dose-response study 5 days following implant. Filter disks treated only with the vehicle for VEGF show no angiogenic response. The siNA were coadminstered with VEGF on a disk in two different siNA concentrations. One concern with the simultaneous administration is that the siNA would not be able to inhibit angiogenesis since VEGF receptors can be stimulated. However, Applicant has observed that in low VEGF doses, the neovascular response reverts to normal, suggesting that the VEGF stimulus is essential for maintaining the angiogenic response. Blocking the production of VEGF receptors using simultaneous administration of anti-VEGF-R mRNA siNA could attenuate the normal neovascularization induced by the filter disk treated with VEGF.

Materials and Methods:

Test Compounds and Controls

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R&D Systems VEGF, carrier free at 75 μ M in 82 mM Tris-Cl, pH 6.9 siNA, 1.67 μ G/ μ L, SITE 2340 (SEQ ID NO: 2; SEQ ID NO: 6) sense/antisense siNA, 1.67 μ G/ μ L, INVERTED CONTROL FOR SITE 2340 (SEQ ID NO: 19; SEQ ID NO: 20) sense/antisense

siNA 1.67 μg/μL, Site 2340 (SEQ ID NO: 419; SEQ ID NO: 420) sense/antisense

Animals

Harlan Sprague-Dawley Rats, Approximately 225-250g
45 males, 5 animals per group.

Husbandry

Animals are housed in groups of two. Feed, water, temperature and humidity are determined according to Pharmacology Testing Facility performance standards (SOP's) which are in accordance with the 1996 Guide for the Care and Use of Laboratory Animals (NRC). Animals are acclimated to the facility for at least 7 days prior to experimentation. During this time, animals are observed for overall health and sentinels will be bled for baseline serology.

Experimental Groups

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Each solution (VEGF and siNAs) was prepared as a 1X solution for final concentrations shown in the experimental groups described in Table III.

siNA Annealing Conditions

1.67mg/mL/strand followed by a 1 hour incubation at 37°C producing 3.34 mg/mL of duplexed siNA. For the 20μg/eye treatment, 6 μLs of the 3.34 mg/mL duplex is injected into the eye (see below). The 3.34 mg/mL duplex siNA can then be serially diluted for dose response assays.

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Preparation of VEGF Filter Disk

For corneal implantation, 0.57 mm diameter nitrocellulose disks, prepared from 0.45 μ m pore diameter nitrocellulose filter membranes (Millipore Corporation), were soaked for 30 min in 1 μ L of 75 μ M VEGF in 82 mM Tris·HCl (pH 6.9) in covered petri dishes on ice. Filter disks soaked only with the vehicle for VEGF (83 mM Tris-Cl pH 6.9) elicit no angiogenic response.

Corneal surgery

The rat corneal model used in this study was a modified from Koch et al. Supra and Pandey et al., supra. Briefly, comeas were irrigated with 0.5% povidone iodine solution followed by normal saline and two drops of 2% lidocaine. Under a dissecting microscope (Leica MZ-6), a stromal pocket was created and a presoaked filter disk (see above) was inserted into the pocket such that its edge was 1 mm from the corneal limbus.

Intraconjunctival injection of test solutions

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Immediately after disk insertion, the tip of a 40-50 µm OD injector (constructed in our laboratory) was inserted within the conjunctival tissue 1 mm away from the edge of the corneal limbus that was directly adjacent to the VEGF-soaked filter disk. Six hundred nanoliters of test solution (siNA, inverted control or sterile water vehicle) were dispensed at a rate of 1.2 µL/min using a syringe pump (Kd Scientific). The injector was then removed, serially rinsed in 70% ethanol and sterile water and immersed in sterile water between each injection. Once the test solution was injected, closure of the eyelid was maintained using microaneurism clips until the animal began to recover gross motor activity. Following treatment, animals were warmed on a heating pad at 37°C.

Quantitation of angiogenic response

administration of 0.4 mg/kg atropine and corneas were digitally imaged. The neovascular surface area (NSA, expressed in pixels) was measured postmortem from blood-filled corneal vessels using computerized morphometry (Image Pro Plus, Media Cybernetics, v2.0). The individual mean NSA was determined in triplicate from three regions of identical size in the area of maximal neovascularization between the filter disk and the limbus. The number of pixels corresponding to the blood-filled corneal vessels in these regions was summated to produce an index of NSA. A group mean NSA was then calculated. Data from each treatment group were normalized to VEGF/siNA vehicle-treated control NSA and finally expressed as percent inhibition of VEGF-induced angiogenesis.

30 Statistics

After determining the normality of treatment group means, group mean percent inhibition of VEGF-induced angiogenesis was subjected to a one-way analysis of variance. This was followed by two post-hoc tests for significance including Dunnett's (comparison to VEGF control) and Tukey-Kramer (all other group mean comparisons) at alpha = 0.05. Statistical analyses were performed using JMP v.3.1.6 (SAS Institute).

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Results are graphically represented in Figure 23. As shown in Figure 23, VEGFR1 site 4229 active siNA at three concentrations were effective at inhibiting angiogenesis compared to the inverted siNA control and the VEGF control. A chemically modified version of the VEGFR1 site 4229 active siNA comprising a sense strand having 2'-deoxy-2'-fluoro pyrimidines and ribo purines with 5' and 3' terminal inverted deoxyabasic residues (SEQ ID NO: 419) and an antisense strand having having 2'-deoxy-2'-fluoro pyrimidines and ribo purines with a terminal 3'-phosphorothioate internucleotide linkage (SEQ ID NO: 420), showed similar inhibition. This result shows siNA molecules of differing chemically modified composition of the invention are capable of significantly inhibiting angiogenesis in vivo.

Example 13: RNAi mediated inhibition of EGFR (HER1) RNA expression

siNA constructs (Table I) were tested for efficacy in reducing EGFR (HER1) RNA expression in A549 cells. A549 cells were plated approximately 24h before transfection in 96-well plates at 5,000-7,500 cells/well, 100 µl/well, such that at the time of transfection cells are 70-90% confluent. For transfection, annealed siNAs were mixed with the transfection reagent (Lipofectamine 2000, Invitrogen) in a volume of 50 µl/well and incubated for 20 min. at room temperature. The siNA transfection mixtures were added to cells to give a final siNA concentration of 25 nM in a volume of 150 µl. Each siNA transfection mixture was added to 3 wells for triplicate siNA treatments. Cells were incubated at 37°C for 24h in the continued presence of the siNA transfection mixture. At 24h, RNA was prepared from each well of treated cells. The supernatants with the transfection mixtures were first removed and discarded, then the cells wre lysed and RNA prepared from each well. Target gene expression following treatment was evaluated by RT-PCR for the target gene and for a control gene (36B4, an RNA polymerase subunit) The triplicate data were averaged and the standard deviations for normalization. determined for each treatment. Normalized data were graphed and the percent reduction

of target mRNA by active siNAs in comparison to their respective inverted control siNAs was determined.

Results of this study are shown in Figure 25. A siNA construct comprising ribonucleotides and 3'-terminal dithymidine caps (RPI#30988/31064) was compared to a chemically modified siNA construct comprising 2'-deoxy-2'-fluoro pyrimidine nucleotides and purine ribonucleotides in which the sense strand of the siNA is further modified with 5' and 3'-terminal inverted deoxyabasic caps and the antisense strand comprises a 3'-terminal phosphorothioate internucleotide linkage (RPI#31300/31301), which was also compared to a matched chemistry inverted control (RPI#31312/31313). In addition, the siNA constructs were also compared to untreated cells, cells transfected with lipid and scrambled siNA constructs (Scram1 and Scram2), and cells transfected with lipid alone (transfection control). As shown in the figure, both siNA constructs significantly reduce EGFR RNA expression. Additional stabilization chemistries as described in Table IV are similarly assayed for activity.

15 Example 14: RNAi mediated inhibition of PKC-alpha RNA expression

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siNA constructs (Table I) are tested for efficacy in reducing PKC-alpha RNA expression in, for example in A549 cells. Cells are plated approximately 24h before transfection in 96-well plates at 5,000-7,500 cells/well, 100 µl/well, such that at the time of transfection cells are 70-90% confluent. For transfection, annealed siNAs are mixed with the transfection reagent (Lipofectamine 2000, Invitrogen) in a volume of 50 µl/well and incubated for 20 min. at room temperature. The siNA transfection mixtures are added to cells to give a final siNA concentration of 25 nM in a volume of 150 µl. Each siNA transfection mixture is added to 3 wells for triplicate siNA treatments. Cells are incubated at 37° for 24h in the continued presence of the siNA transfection mixture. At 24h, RNA is prepared from each well of treated cells. The supernatants with the transfection mixtures are first removed and discarded, then the cells are lysed and RNA prepared from each well. Target gene expression following treatment is evaluated by RT-PCR for the target gene and for a control gene (36B4, an RNA polymerase subunit) for normalization. The triplicate data is averaged and the standard deviations determined for each treatment. Normalized data are graphed and the percent reduction of target mRNA by active siNAs in comparison to their respective inverted control siNAs was determined.

In a non-limiting example, siNA constructs were screened for activity (see Figure 26) and compared to untreated cells, scrambled siNA control constructs (Scram1 and Scram2), and cells transfected with lipid alone (transfection control). As shown in Figure 26, the siNA constructs significantly reduce PKC-alpha RNA expression. Leads generated from such a screen are then further assayed. In a non-limiting example, siNA constructs comprising ribonucleotides and 3'-terminal dithymidine caps are assayed along with a chemically modified siNA construct comprising 2'-deoxy-2'-fluoro pyrimidine nucleotides and purine ribonucleotides, in which the sense strand of the siNA is further modified with 5' and 3'-terminal inverted deoxyabasic caps and the antisense strand comprises a 3'-terminal phosphorothioate internucleotide linkage. Additional stabilization chemistries as described in Table IV are similarly assayed for activity. These siNA constructs are compared to appropriate matched chemistry inverted controls. In addition, the siNA constructs are also compared to untreated cells, cells transfected with lipid and scrambled siNA constructs, and cells transfected with lipid alone (transfection control).

Example 15: RNAi mediated inhibition of Myc RNA expression

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siNA constructs (Table I) were tested for efficacy in reducing Myc (c-Myc) RNA expression in 293T cells. 293T cells were plated approximately 24h before transfection in 96-well plates at 5,000-7,500 cells/well, 100 µl/well, such that at the time of transfection cells were 70-90% confluent. For transfection, annealed siNAs were mixed with the transfection reagent (Lipofectamine 2000, Invitrogen) in a volume of 50 µl/well and incubated for 20 min. at room temperature. The siNA transfection mixtures were added to cells to give a final siNA concentration of 25 nM in a volume of 150 µl. Each siNA transfection mixture was added to 3 wells for triplicate siNA treatments. Cells were incubated at 37°C for 24h in the continued presence of the siNA transfection mixture. At 24h, RNA was prepared from each well of treated cells. The supernatants with the transfection mixtures were first removed and discarded, then the cells were lysed and RNA prepared from each well. Target gene expression following treatment was evaluated by RT-PCR for the target gene and for a control gene (36B4, an RNA polymerase subunit) for normalization. The triplicate data were averaged and the standard deviations determined for each treatment. Normalized data were graphed and

the percent reduction of target mRNA by active siNAs in comparison to their respective inverted control siNAs was determined.

Results of this study are shown in Figure 27. A screen of siNA constructs was compared to untreated cells, scrambled siNA control constructs (Scram1 and Scram2), and cells transfected with lipid alone (transfection control). As shown in the figure, three of the siNA constructs (RPI 30993/31069; RPI 30995/31071; and RPI 30996/31072) significantly reduce c-Myc RNA expression. Additional stabilization chemistries as described in Table IV are similarly assayed for activity.

Example 16: RNAi mediated inhibition of BCL2 RNA expression

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siNA constructs (Table I) are tested for efficacy in reducing BCL2 RNA expression in, for example, A549 cells. Cells are plated approximately 24h before transfection in 96well plates at 5,000-7,500 cells/well, 100 μl/well, such that at the time of transfection cells are 70-90% confluent. For transfection, annealed siNAs are mixed with the transfection reagent (Lipofectamine 2000, Invitrogen) in a volume of 50 µl/well and incubated for 20 min. at room temperature. The siNA transfection mixtures are added to cells to give a final siNA concentration of 25 nM in a volume of 150 µl. Each siNA transfection mixture is added to 3 wells for triplicate siNA treatments. Cells are incubated at 37° for 24h in the continued presence of the siNA transfection mixture. At 24h, RNA is prepared from each well of treated cells. The supernatants with the transfection mixtures are first removed and discarded, then the cells are lysed and RNA prepared from each well. Target gene expression following treatment is evaluated by RT-PCR for the target gene and for a control gene (36B4, an RNA polymerase subunit) for normalization. The triplicate data is averaged and the standard deviations determined for each treatment. Normalized data are graphed and the percent reduction of target mRNA by active siNAs in comparison to their respective inverted control siNAs is determined.

In a non-limiting example, A549 cells were transfected with 0.25 ug/well of lipid complexed with 25 nM siNA. A siNA construct comprising ribonucleotides and 3'-terminal dithymidine caps (RPI#30998/31074) was tested along with a chemically modified siNA construct comprising 2'-deoxy-2'-fluoro pyrimidine nucleotides and purine ribonucleotides in which the sense strand of the siNA is further modified with 5' and 3'-terminal inverted deoxyabasic caps and the antisense strand comprises a 3'-

terminal phosphorothioate internucleotide linkage (RPI#31368/31369), which was also compared to a matched chemistry inverted control (RPI#31370/31371) and a chemically modified siNA construct comprising 2'-deoxy-2'-fluoro pyrimidine and 2'-deoxy-2'-fluoro purine nucleotides in which the sense strand of the siNA is further modified with 5' and 3'-terminal inverted deoxyabasic caps and the antisense strand comprises a 3'-terminal phosphorothioate internucleotide linkage (RPI#31372/31373) which was also compared to a matched chemistry inverted control (RPI#31374/31375). In addition, the siNA constructs were also compared to untreated cells, cells transfected with lipid and scrambled siNA constructs (Scram1 and Scram2), and cells transfected with lipid alone (transfection control). As shown in Figure 28, the siNA constructs significantly reduce BCL2 RNA expression compared to scrambled, untreated, and transfection controls. Additional stabilization chemistries as described in Table IV are similarly assayed for activity.

Example 17: RNAi mediated inhibition of CHK-1 RNA expression

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siNA constructs (Table I) were tested for efficacy in reducing CHK-1 RNA expression in A549 cells. A549 cells were plated approximately 24h before transfection in 96-well plates at 5,000-7,500 cells/well, 100 µl/well, such that at the time of transfection cells are 70-90% confluent. For transfection, annealed siNAs were mixed with the transfection reagent (Lipofectamine 2000, Invitrogen) in a volume of 50 µl/well and incubated for 20 min. at room temperature. The siNA transfection mixtures were added to cells to give a final siNA concentration of 25 nM in a volume of 150 µl. Each siNA transfection mixture was added to 3 wells for triplicate siNA treatments. Cells were incubated at 37° for 24h in the continued presence of the siNA transfection mixture. At 24h, RNA was prepared from each well of treated cells. The supernatants with the transfection mixtures were first removed and discarded, then the cells were lysed and RNA prepared from each well. Target gene expression following treatment was evaluated by RT-PCR for the target gene and for a control gene (36B4, an RNA polymerase subunit) for normalization. The triplicate data were averaged and the standard deviations determined for each treatment. Normalized data were graphed and the percent reduction of target mRNA by active siNAs in comparison to their respective inverted control siNAs was determined.

Results of this study are shown in Figure 29. A siNA construct comprising ribonucleotides and 3'-terminal dithymidine caps (RPI#31003/31079) and a chemically modified siNA construct comprising 2'-deoxy-2'-fluoro pyrimidine nucleotides and purine ribonucleotides in which the sense strand of the siNA is further modified with 5' and 3'-terminal inverted deoxyabasic caps and in which the antisense strand comprises a 3'-terminal phosphorothioate internucleotide linkage (RPI#31302/31303), were compared to a matched chemistry inverted control (RPI#31314/31325). In addition, the siNA constructs were also compared to untreated cells, cells transfected with lipid and scrambled siNA constructs (Scram1 and Scram2), and cells transfected with lipid alone (transfection control). As shown in the figure, both siNA constructs significantly reduce CHK-1 RNA expression compared to appropriate controls. Additional stabilization chemistries as described in Table IV are similarly assayed for activity.

Example 18: RNAi mediated inhibition of BACE RNA expression

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siNA constructs (Table I) are tested for efficacy in reducing BACE RNA expression in, for example in A549 cells. Cells are plated approximately 24h before transfection in 96well plates at 5,000-7,500 cells/well, 100 μl/well, such that at the time of transfection cells are 70-90% confluent. For transfection, annealed siNAs are mixed with the transfection reagent (Lipofectamine 2000, Invitrogen) in a volume of 50 µl/well and incubated for 20 min. at room temperature. The siNA transfection mixtures are added to cells to give a final siNA concentration of 25 nM in a volume of 150 µl. Each siNA transfection mixture is added to 3 wells for triplicate siNA treatments. Cells are incubated at 37°C for 24h in the continued presence of the siNA transfection mixture. At 24h, RNA is prepared from each well of treated cells. The supernatants with the transfection mixtures are first removed and discarded, then the cells are lysed and RNA prepared from each well. Target gene expression following treatment is evaluated by RT-PCR for the target gene and for a control gene (36B4, an RNA polymerase subunit) for normalization. The triplicate data is averaged and the standard deviations determined for each treatment. Normalized data are graphed and the percent reduction of target mRNA by active siNAs in comparison to their respective inverted control siNAs was determined.

In a non-limiting example, siNA constructs were screened for activity (see Figure 30) and compared to untreated cells, scrambled siNA control constructs (Scram1 and

Scram2), and cells transfected with lipid alone (transfection control). As shown in Figure 30, the siNA constructs significantly reduce BACE RNA expression. Leads generated from such a screen are then further assayed. In a non-limiting example, siNA constructs comprising ribonucleotides and 3'-terminal dithymidine caps are assayed along with a chemically modified siNA construct comprising 2'-deoxy-2'-fluoro pyrimidine nucleotides and purine ribonucleotides, in which the sense strand of the siNA is further modified with 5' and 3'-terminal inverted deoxyabasic caps and the antisense strand comprises a 3'-terminal phosphorothioate internucleotide linkage. Additional stabilization chemistries as described in Table IV are similarly assayed for activity. These siNA constructs are compared to appropriate matched chemistry inverted controls. In addition, the siNA constructs are also compared to untreated cells, cells transfected with lipid and scrambled siNA constructs, and cells transfected with lipid alone (transfection control).

Example 19: RNAi mediated inhibition of cyclin D1 RNA expression

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siNA constructs (Table I) were tested for efficacy in reducing cyclin D1 RNA expression in A549 cells. A549 cells were plated approximately 24h before transfection in 96-well plates at 5,000-7,500 cells/well, 100 µl/well, such that at the time of transfection cells are 70-90% confluent. For transfection, annealed siNAs were mixed with the transfection reagent (Lipofectamine 2000, Invitrogen) in a volume of 50 μ l/well and incubated for 20 min. at room temperature. The siNA transfection mixtures were added to cells to give a final siNA concentration of 25 nM in a volume of 150 µl. Each siNA transfection mixture was added to 3 wells for triplicate siNA treatments. Cells were incubated at 37° for 24h in the continued presence of the siNA transfection mixture. At 24h, RNA was prepared from each well of treated cells. The supernatants with the transfection mixtures were first removed and discarded, then the cells were lysed and RNA prepared from each well. Target gene expression following treatment was evaluated by RT-PCR for the target gene and for a control gene (36B4, an RNA polymerase subunit) for normalization. The triplicate data were averaged and the standard deviations determined for each treatment. Normalized data were graphed and the percent reduction of target mRNA by active siNAs in comparison to their respective inverted control siNAs was determined.

Results of this study are shown in Figure 31. A siNA construct comprising ribonucleotides and 3'-terminal dithymidine caps (RPI#30988/31064) was assayed along with a chemically modified siNA construct comprising 2'-deoxy-2'-fluoro pyrimidine nucleotides and purine ribonucleotides in which the sense strand of the siNA is further modified with 5' and 3'-terminal inverted deoxyabasic caps and the antisense strand comprises a 3'-terminal phosphorothioate internucleotide linkage (RPI#31300/3130), which was also compared to a matched chemistry inverted control (RPI#31312/31313). In addition, the siNA constructs were also compared to untreated cells, cells transfected with lipid and scrambled siNA constructs (Scram1 and Scram2), and cells transfected with lipid alone (transfection control). As shown in the figure, both siNA constructs significantly reduce cyclin D1 RNA expression. Additional stabilization chemistries as described in Table IV are similarly assayed for activity.

Example 20: RNAi mediated inhibition of PTP-1B RNA expression

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siNA constructs (Table I) were tested for efficacy in reducing PTP-1B RNA expression in A549 cells. A549 cells were plated approximately 24h before transfection in 96-well plates at 5,000-7,500 cells/well, 100 µl/well, such that at the time of transfection cells are 70-90% confluent. For transfection, annealed siNAs were mixed with the transfection reagent (Lipofectamine 2000, Invitrogen) in a volume of 50 µl/well and incubated for 20 min. at room temperature. The siNA transfection mixtures were added to cells to give a final siNA concentration of 25 nM in a volume of 150 μ l. Each siNA transfection mixture was added to 3 wells for triplicate siNA treatments. Cells were incubated at 37° for 24h in the continued presence of the siNA transfection mixture. At 24h, RNA was prepared from each well of treated cells. The supernatants with the transfection mixtures were first removed and discarded, then the cells were lysed and RNA prepared from each well. Target gene expression following treatment was evaluated by RT-PCR for the target gene and for a control gene (36B4, an RNA polymerase subunit) for normalization. The triplicate data were averaged and the standard deviations determined for each treatment. Normalized data were graphed and the percent reduction of target mRNA by active siNAs in comparison to their respective inverted control siNAs was determined.

Results of this study are shown in Figure 32. A siNA construct comprising ribonucleotides and 3'-terminal dithymidine caps (RPI#31018/31094) was assayed along with a chemically modified siNA construct comprising 2'-deoxy-2'-fluoro pyrimidine nucleotides and purine ribonucleotides in which the sense strand of the siNA is further modified with 5' and 3'-terminal inverted deoxyabasic caps and the antisense strand comprises a 3'-terminal phosphorothioate internucleotide linkage (RPI#31306/31307), which was also compared to a matched chemistry inverted control (RPI#31318/31319). In addition, the siNA constructs were also compared to untreated cells, cells transfected with lipid and scrambled siNA constructs (Scram1 and Scram2), and cells transfected with lipid alone (transfection control). As shown in the figure, both siNA constructs significantly reduce PTP-1B RNA expression. Additional stabilization chemistries as described in Table IV are similarly assayed for activity.

Example 21: RNAi mediated inhibition of ERG2 RNA expression

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siNA constructs (Table I) are tested for efficacy in reducing ERG2 RNA expression in, for example in DLD1 cells. Cells are plated approximately 24h before transfection in 96-well plates at 5,000-7,500 cells/well, 100 µl/well, such that at the time of transfection cells are 70-90% confluent. For transfection, annealed siNAs are mixed with the transfection reagent (Lipofectamine 2000, Invitrogen) in a volume of 50 μ l/well and incubated for 20 min. at room temperature. The siNA transfection mixtures are added to cells to give a final siNA concentration of 25 nM in a volume of 150 μ l. Each siNA transfection mixture is added to 3 wells for triplicate siNA treatments. Cells are incubated at 37° for 24h in the continued presence of the siNA transfection mixture. At 24h, RNA is prepared from each well of treated cells. The supernatants with the transfection mixtures are first removed and discarded, then the cells are lysed and RNA prepared from each well. Target gene expression following treatment is evaluated by RT-PCR for the target gene and for a control gene (36B4, an RNA polymerase subunit) for normalization. The triplicate data is averaged and the standard deviations determined for each treatment. Normalized data are graphed and the percent reduction of target mRNA by active siNAs in comparison to their respective inverted control siNAs was determined.

In a non-limiting example, siNA constructs were screened for activity (see Figure 33) and compared to untreated cells, scrambled siNA control constructs (Scram1 and

Scram2), and cells transfected with lipid alone (transfection control). As shown in Figure 33, the siNA constructs significantly reduce of ERG2 RNA expression. Leads generated from such a screen are then further assayed. In a non-limiting example, siNA constructs comprising ribonucleotides and 3'-terminal dithymidine caps are assayed along with a chemically modified siNA construct comprising 2'-deoxy-2'-fluoro pyrimidine nucleotides and purine ribonucleotides, in which the sense strand of the siNA is further modified with 5' and 3'-terminal inverted deoxyabasic caps and the antisense strand comprises a 3'-terminal phosphorothioate internucleotide linkage. Additional stabilization chemistries as described in Table IV are similarly assayed for activity. These siNA constructs are compared to appropriate matched chemistry inverted controls. In addition, the siNA constructs are also compared to untreated cells, cells transfected with lipid and scrambled siNA constructs, and cells transfected with lipid alone (transfection control). Additional stabilization chemistries as described in Table IV are similarly assayed for activity.

15 Example 22: RNAi mediated inhibition of PCNA RNA expression

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siNA constructs (Table I) were tested for efficacy in reducing PCNA RNA expression in A549 cells. A549 cells were plated approximately 24h before transfection in 96-well plates at 5,000-7,500 cells/well, 100 µl/well, such that at the time of transfection cells are 70-90% confluent. For transfection, annealed siNAs were mixed with the transfection reagent (Lipofectamine 2000, Invitrogen) in a volume of 50 µl/well and incubated for 20 min. at room temperature. The siNA transfection mixtures were added to cells to give a final siNA concentration of 25 nM in a volume of 150 µl. Each siNA transfection mixture was added to 3 wells for triplicate siNA treatments. Cells were incubated at 37° for 24h in the continued presence of the siNA transfection mixture. At 24h, RNA was prepared from each well of treated cells. The supernatants with the transfection mixtures were first removed and discarded, then the cells were lysed and RNA prepared from each well. Target gene expression following treatment was evaluated by RT-PCR for the target gene and for a control gene (36B4, an RNA polymerase subunit) for normalization. The triplicate data were averaged and the standard deviations determined for each treatment. Normalized data were graphed and the percent reduction of target mRNA by active siNAs in comparison to their respective inverted control siNAs was determined.

Results of this study are shown in Figure 34. A siNA construct comprising ribonucleotides and 3'-terminal dithymidine caps (RPI#31035/31111) was assayed along with a chemically modified siNA construct comprising 2'-deoxy-2'-fluoro pyrimidine nucleotides and purine ribonucleotides in which the sense strand of the siNA is further modified with 5' and 3'-terminal inverted deoxyabasic caps and the antisense strand comprises a 3'-terminal phosphorothioate internucleotide linkage (RPI#31310/31311), which was also compared to a matched chemistry inverted control (RPI#31322/31323). In addition, the siNA constructs were also compared to untreated cells, cells transfected with lipid and scrambled siNA constructs (Scram1 and Scram2), and cells transfected with lipid alone (transfection control). As shown in the figure, both siNA constructs significant reduce PCNA RNA expression. Additional stabilization chemistries as described in Table IV are similarly assayed for activity.

Example 23: Indications

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The siNA molecules of the invention can be used to treat a variety of diseases and conditions through modulation of gene expression. Using the methods described herein, chemically modified siNA molecules can be designed to modulate the expression any number of target genes, including but not limited to genes associated with cancer, metabolic diseases, infectious diseases such as viral, bacterial or fungal infections, neurologic diseases, musculoskeletal diseases, diseases of the immune system, diseases associated with signaling pathways and cellular messengers, and diseases associated with transport systems including molecular pumps and channels.

Non-limiting examples of various viral genes that can be targeted using siRNA molecules of the invention include Hepatitis C Virus (HCV, for example Genbank Accession Nos: D11168, D50483.1, L38318 and S82227), Hepatitis B Virus (HBV, for example GenBank Accession No. AF100308.1), Human Immunodeficiency Virus type 1 (HIV-1, for example GenBank Accession No. U51188), Human Immunodeficiency Virus type 2 (HIV-2, for example GenBank Accession No. X60667), West Nile Virus (WNV for example GenBank accession No. NC_001563), cytomegalovirus (CMV for example GenBank Accession No. NC_001347), respiratory syncytial virus (RSV for example GenBank Accession No. NC_001781), influenza virus (for example example GenBank Accession No. AF037412, rhinovirus (for example, GenBank accession numbers:

D00239, X02316, X01087, L24917, M16248, K02121, X01087), papillomavirus (for example GenBank Accession No. NC_001353), Herpes Simplex Virus (HSV for example GenBank Accession No. NC_001345), and other viruses such as HTLV (for example GenBank Accession No. AJ430458). Due to the high sequence variability of many viral genomes, selection of siRNA molecules for broad therapeutic applications would likely involve the conserved regions of the viral genome. Nonlimiting examples of conserved regions of the viral genomes include but are not limited to 5'-Non Coding Regions (NCR), 3'- Non Coding Regions (NCR) and/or internal ribosome entry sites (IRES). siRNA molecules designed against conserved regions of various viral genomes will enable efficient inhibition of viral replication in diverse patient populations and may ensure the effectiveness of the siRNA molecules against viral quasi species which evolve due to mutations in the non-conserved regions of the viral genome.

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Non-limiting examples of human genes that can be targeted using siRNA molecules of the invention using methods described herein include any human RNA sequence, for example those commonly referred to by Genbank Accession Number. These RNA sequences can be used to design siRNA molecules that inhibit gene expression and therefore abrogate diseases, conditions, or infections associated with expression of those genes. Such non-limiting examples of human genes that can be targeted using siRNA molecules of the invention include VEGFr (VEGFr-1 for example GenBank Accession No. XM 067723, VEGFr-2 for example GenBank Accession No. AF063658), HER1, HER2, HER3, and HER4 (for example Genbank Accession Nos: NM_005228, NM 004448, NM 001982, and NM 005235 respectively), telomerase (TERT, for example GenBank Accession No. NM 003219), telomerase RNA (for example GenBank Accession No. U86046), NFkappaB, Rel-A (for example GenBank Accession No. NM 005228), NOGO (for example GenBank Accession No. AB020693), NOGOr (for example GenBank Accession No. XM 015620), RAS (for example GenBank Accession No. NM 004283), RAF (for example GenBank Accession No. XM 033884), CD20 (for example GenBank Accession No. X07203), METAP2 (for example GenBank Accession No. NM_003219), CLCA1 (for example GenBank Accession No. NM_001285), phospholamban (for example GenBank Accession No. NM 002667), PTP1B (for example GenBank Accession No. M31724), and others, for example, those shown in Table III.

The siNA molecule of the invention can also be used in a variety of agricultural applications involving modulation of endogenous or exogenous gene expression in plants using siNA, including use as insecticidal, antiviral and anti-fungal agents or modulate plant traits such as oil and starch profiles and stress resistance.

5 Example 24: Diagnostic uses

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The siNA molecules of the invention can be used in a variety of diagnostic applications, such as in the identification of molecular targets (e.g., RNA) in a variety of applications, for example, in clinical, industrial, environmental, agricultural and/or research settings. Such diagnostic use of siNA molecules involves utilizing reconstituted RNAi systems, for example, using cellular lysates or partially purified cellular lysates. siNA molecules of this invention can be used as diagnostic tools to examine genetic drift and mutations within diseased cells or to detect the presence of endogenous or exogenous, for example viral, RNA in a cell. The close relationship between siNA activity and the structure of the target RNA allows the detection of mutations in any region of the molecule, which alters the base-pairing and three-dimensional structure of the target RNA. By using multiple siNA molecules described in this invention, one can map nucleotide changes, which are important to RNA structure and function in vitro, as well as in cells and tissues. Cleavage of target RNAs with siNA molecules can be used to inhibit gene expression and define the role of specified gene products in the progression of disease or infection. In this manner, other genetic targets can be defined as important mediators of the disease. These experiments will lead to better treatment of the disease progression by affording the possibility of combination therapies (e.g., multiple siNA molecules targeted to different genes, siNA molecules coupled with known small molecule inhibitors, or intermittent treatment with combinations siNA molecules and/or other chemical or biological molecules). Other in vitro uses of siNA molecules of this invention are well known in the art, and include detection of the presence of mRNAs associated with a disease, infection, or related condition. Such RNA is detected by determining the presence of a cleavage product after treatment with a siNA using standard methodologies, for example, fluorescence resonance emission transfer (FRET).

In a specific example, siNA molecules that cleave only wild-type or mutant forms of the target RNA are used for the assay. The first siNA molecules (i.e., those that cleave

only wild-type forms of target RNA) are used to identify wild-type RNA present in the sample and the second siNA molecules (i.e., those that cleave only mutant forms of target RNA) are used to identify mutant RNA in the sample. As reaction controls, synthetic substrates of both wild-type and mutant RNA are cleaved by both siNA molecules to demonstrate the relative siNA efficiencies in the reactions and the absence of cleavage of the "non-targeted" RNA species. The cleavage products from the synthetic substrates also serve to generate size markers for the analysis of wild-type and mutant RNAs in the sample population. Thus, each analysis requires two siNA molecules, two substrates and one unknown sample, which is combined into six reactions. The presence of cleavage products is determined using an RNase protection assay so that full-length and cleavage fragments of each RNA can be analyzed in one lane of a polyacrylamide gel. It is not absolutely required to quantify the results to gain insight into the expression of mutant RNAs and putative risk of the desired phenotypic changes in target cells. The expression of mRNA whose protein product is implicated in the development of the phenotype (i.e., disease related or infection related) is adequate to establish risk. If probes of comparable specific activity are used for both transcripts, then a qualitative comparison of RNA levels is adequate and decreases the cost of the initial diagnosis. Higher mutant form to wildtype ratios are correlated with higher risk whether RNA levels are compared qualitatively or quantitatively.

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All patents and publications mentioned in the specification are indicative of the levels of skill of those skilled in the art to which the invention pertains. All references cited in this disclosure are incorporated by reference to the same extent as if each reference had been incorporated by reference in its entirety individually.

One skilled in the art would readily appreciate that the present invention is well adapted to carry out the objects and obtain the ends and advantages mentioned, as well as those inherent therein. The methods and compositions described herein as presently representative of preferred embodiments are exemplary and are not intended as limitations on the scope of the invention. Changes therein and other uses will occur to those skilled in the art, which are encompassed within the spirit of the invention, are defined by the scope of the claims.

It will be readily apparent to one skilled in the art that varying substitutions and modifications can be made to the invention disclosed herein without departing from the scope and spirit of the invention. Thus, such additional embodiments are within the scope of the present invention and the following claims. The present invention teaches one skilled in the art to test various combinations and/or substitutions of chemical modifications described herein toward generating nucleic acid constructs with improved activity for mediating RNAi activity. Such improved activity can comprise improved stability, improved bioavailability, and/or improved activation of cellular responses mediating RNAi. Therefore, the specific embodiments described herein are not limiting and one skilled in the art can readily appreciate that specific combinations of the modifications described herein can be tested without undue experimentation toward identifying siNA molecules with improved RNAi activity.

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The invention illustratively described herein suitably can be practiced in the absence of any element or elements, limitation or limitations that are not specifically disclosed herein. Thus, for example, in each instance herein any of the terms "comprising", "consisting essentially of", and "consisting of" may be replaced with either of the other two terms. The terms and expressions which have been employed are used as terms of description and not of limitation, and there is no intention that in the use of such terms and expressions of excluding any equivalents of the features shown and described or portions thereof, but it is recognized that various modifications are possible within the scope of the invention claimed. Thus, it should be understood that although the present invention has been specifically disclosed by preferred embodiments, optional features, modification and variation of the concepts herein disclosed may be resorted to by those skilled in the art, and that such modifications and variations are considered to be within the scope of this invention as defined by the description and the appended claims.

In addition, where features or aspects of the invention are described in terms of Markush groups or other grouping of alternatives, those skilled in the art will recognize that the invention is also thereby described in terms of any individual member or subgroup of members of the Markush group or other group.

Table

Seq!D	186	187	188	189	190	191	192	193	194	195	196	197	198	199	200	201	202	203	204	205	900
Sequence	B uuccuccuGGAAAuucAAcTT B	B ccucucAuGAuGcuGGuGuTT B	B cGAuAGcuGAAAAcAuucGTT B	B AAuGcAGcuGAuGAAuccATT B	GuuGAAuuuccAGGAGGAATsT	AcAccAGcAucAuGAGAGGTsT	cGAAuGuuucAGcuAucGTsT	uGGAuucAucAGcuGcAuuTsT	UUCCUCCUGGAAAUUCAACTT	CCUCUCAUGAUGCUGGUGUTT	CGAUAGCUGAAAACAUUCGTT	AAUGCAGCUGAUGAAUCCATT	GUUGAAUUUCCAGGAGGAATT	ACACCAGCAUCAUGAGAGGTT	CGAAUGUUUCAGCUAUCGTT	UGGAUUCAUCAGCUGCAUUTT	B uucGAGAAGGucAucAGcATT B	B ccAGGuGucuAGAGGcAAcTT B	B AccAAGcuuAAGGAGAGGATT B	B cGGuuGAccuucuGAAcAuTT B	1 1 1 1 1 1 1 1 1 1 1 1 1 1 1 1 1 1 1 1
# Aliases	17 ABCB1:120U21 siRNA stab04	30938 ABCB1:620U21 siRNA stab04	30939 ABCB1:1869U21 siRNA stab04	30940 ABCB1:2336U21 siRNA stab04	11 ABCB1:138L21 siRNA (120C) stab05	30942 ABCB1:638L21 siRNA (620C) stab05	30943 ABCB1:1887L21 siRNA (1869C) stab05	4 ABCB1:2354L21 siRNA (2336C) stab05	_	31014 ABCB1:620U21 siRNA	31015 ABCB1:1869U21 siRNA	31016 ABCB1:2336U21 sIRNA	31089 ABCB1:138L21 siRNA (120C)	10 ABCB1:638L21 siRNA (620C)	11 ABCB1:1887L21 siRNA (1869C)	antisense 31092 ABCB1:2354L21 siRNA (2336C)	30721 ADORA1:921U21 siRNA stab04		23 ADORA1:1821U21 siRNA stab04	30724 ADORA1:2775U21 siRNA stab04	
RP#	30937	3093	3093	3094	30941			30944	31013	3101	3101	3101	3108	31090	31091	3109	3072	30722	30723	3072	
strand	seuse	sense	seuse	sense	antisense	antisense	antisense	antisense	seuse	seuse	seuse	seuse	antisense	antisense	antisense	antisense	seuse	seuse	seuse	seuse	
Seq	-	2	က	4	1	2	က	4	1	2	3	4	-	7	က	4	က	9	7	ω	
Target Sequence	CAUUCCUCCUGGAAAUUCAACCU	UUCCUCAUGAUGCUGGUGUUU	CACGAUAGCUGAAAACAUUCGCU	AAAAUGCAGCUGAUGAAUCCAAA	CAUUCCUCCUGGAAAUUCAACCU	UNCCUCUCAUGAUGCUGGUGUUU	CACGAUAGCUGAAAACAUUCGCU	AAAAUGCAGCUGAUGAAUCCAAA	CAUUCCUCCUGGAAAUUCAACCU	UNCCUCUCAUGAUGCUGGUGUUU	CACGAUAGCUGAAAACAUUCGCU		CAUUCCUCCUGGAAAUUCAACCU	UUCCUCUCAUGAUGCUGGUGUUU	CACGAUAGCUGAAAACAUUCGCU	AAAAUGCAGCUGAUGAAUCCAAA	AGUUCGAGAGGUCAUCAGCAUG	GACCAGGUGUCUAGAGGCAACAG	GGACCAAGCUUAAGGAGAGGAGA	GUCGGUUGACCUUCUGAACAUGA	0.100.00.00.00.00.00.00.00.00.00.00.00.0
Targe t Pos	118	618	1867	2334	118	618	1867	2334	118	618	1867	2334	118	618	1867	2334	919	1621	1819	2773	9
Target Targe	ABCB1	ABCB1	ABCB1	ABCB1	ABCB1	ABCB1	ABCB1	ABCB1	ABCB1	ABCB1	ABCB1	ABCB1	ABCB1	ABCB1	ABCB1	ABCB1	ADORA 1	ADORA 1621	ADORA 1819	ADORA 1	

	207	208	209	210	211	212	213	214	215	216	217	218	219	220	221	222	223	224	225	226	227	228	229	230
	GuuGccucuAGAcAccuGGTsT	uccucuuAAGcuuGGuTsT	AuGuucAGAAGGucAAccGTsT	UUCGAGAAGGUCAUCAGCATT	CCAGGUGUCUAGAGGCAACTT	ACCAAGCUUAAGGAGGAGTT	CGGUUGACCUUCUGAACAUTT	UGCUGAUGACCUUCUCGAATT	GUUGCCUCUAGACACCUGGTT	UCCUCUCCUUAAGCUUGGUTT	AUGUUCAGAAGGUCAACCGTT	ACCAUCAAUAAGGAAGAAGTT	AUCAAUAAGGAAGAAGCCCTT	GACCAUCAAUAAGGAAGAATT	AUAAGGAAGAAGCCCUUCATT	CUUCUUCCUUAUUGAUGGUTT	GGCCUCCUCCUCAUGAUTT	UCCUUCCUUAUUGAUGGUCTT	UGAAGGGCUUCUUCCUUAUTT	GAUUUAAGCAGAGUUCAAATT	AGAGUUCAAAAGCCCUUCATT	CAGAGUUCAAAAGCCCUUCTT	AUUUAAGCAGAGUUCAAAATT	UUUGAACUCUGCUUAAAUCTT
(921C) stab05		ADORA1:1839L21 siRNA (1821C) stab05	antisense 30728 ADORA1:2793L21 siRNA (2775C) stab05	31041 ADORA1:921U21 siRNA	31042 ADORA1:1623U21 siRNA	31043 ADORA1:1821U21 siRNA	31044 ADORA1:2775U21 siRNA	31117 ADORA1:939L21 siRNA (921C)	31118 ADORA1:1641L21 siRNA (1623C)	31119 ADORA1:1839L21 siRNA (1821C)	31120 ADORA1:2793L21 sIRNA (2775C)	31594 b2a2:283U21 siRNA	31595 b2a2:286U21 siRNA	31596 b2a2:282U21 siRNA	31597 b2a2:290U21 siRNA	31598 b2a2:301L21 siRNA (283C)	31599 b2a2:304L21 siRNA (286C)	31600 b2a2:300L21 siRNA (282C)	31601 b2a2:308L21 siRNA (290C)	b3a2:356U21 siRNA	31603 b3a2:365U21 siRNA	31604 b3a2:364U21 siRNA	31605 b3a2:357U21 siRNA	31606 b3a2:374L21 siRNA (356C)
	30726	30727	30728	31041	31042	31043	31044	31117			31120	31594	31595	31596	31597	31598	31599	31600		31602	31603	31604	31605	31606
	antisense	antisense	antisense	seuse	sense	sense	sense	antisense	antisense	antisense	antisense	esues	esues	sense	esues	antisense	antisense	antisense	antisense	sense	seuse	seuse	seuse	antisense
	9	7	8	ည	9	7	8	5	9	2	8	6	10	11	12	တ	10	11	12	13	14	15	16	13
_		GGACCAAGCUUAAG	4.	AGUUCGAGAGGUCAUCAGCAUG	GACCAGGUGUCUAGAGGCAACAG	GGACCAAGCUUAAGGAGAGGAGA	GUCGGUUGACCUUCUGAACAUGA					-	_4		CAAUAAGGAAGAAGCCCUUCAGC	UGACCAUCAAUAAGGAAGAGCC	CCAUCAAUAAGGAAGAAGCCCUU	CUGACCAUCAAUAAGGAAGAAGC			_	AGCAGAGUUCAAAAGCCCUUCAG	GGAUUUAAGCAGAGUUCAAAAGC	UGGAUUUAAGCAGAGUUCAAAAG
				919	1621	1819	2773				٠, ا	283	286	282	230	301	304	300	308	356	365	364	357	374
-	ADORA 1	ADORA 1	ADORA 1	ADORA 1	ADORA 1	ADORA 1	ADORA 1	ADORA 1	ADORA 1	ADORA 1	ADORA 1	b2a2	b2a2	b2a2	b2a2	b2a2	b2a2	b2a2	b2a2	b3a2	p3a2	b3a2	b3a2	p3a2

b3a2	383	GCAGAGUUCAAAAGCCCUUCAGC	14	antisense		31607 b3a2:383L21 siRNA	UGAAGGGCUUUUGAACUCUTT	234
b3a2	382	AGCAGAGUUCAAAAGCCCUUCAG	15	antisense	31608		GAAGGGCUUUUGAACUCUGTT	232
b3a2	375	GGAUUUAAGCAGAGUUCAAAAGC	16	antisansa		31609 h322.3751 21 cibnia		
BACE			2	DCIIIO CIIII D	91009	(357C)	UUUUGAACUCUGCUUAAAUTT	233
DACE		AAUGGGUGAGGUUACCAACCAGU	17	sense	30729	BACE:1492U21 siRNA stab04	B uGGGuGAGGuuAccAAccATT B	234
BACE			18	seuse	30730	30730 BACE:1755U21 siRNA	B AccuuGGAcAuGGAAGAcuTT B	235
BACE	$\overline{}$		19	sense	30732	BACE:3585U21 siRNA	B uGGGAccuGcuAAGuGuGGTT B	236
BACE		$\overline{}$	17	antisense	30733	BACE:1510L21 siRNA (1492C) stab05	uGGuuGGuAAccucAcccATsT	237
BACE	1753	UCACCUUGGACAUGGAAGACUGU	18	antisense		30734 BACE:1773L21 siRNA	AGucuuccAuGuccAAGGuTsT	238
BACE	3583	UAUGGGACCUGCUAAGUGUGGAA	19	antisense	30736	30736 BACE:3603L21 siRNA	ccAcAcuuAGcAGGucccATsT	239
BACE	_	_	17	sense	31005	31005 BACE:14921121 siRNA		
BACE	1753	UCACCUUGGACAUGGAAGACUGU	18	sense	31006	31006 BACE:1755U21 siRNA	ACCILIGGACALIGGAACITT	240
DACE DACE	_		20	sense	31007	31007 BACE:2459U21 siRNA	UAACAUUGGUGCAAAGAUUTT	242
מ ללם	-		19	sense	31008	31008 BACE:3585U21 siRNA	UGGGACCUGCUAAGUGUGGTT	243
DACE		AAUGGGUGAGGUUACCAACCAGU	17	antisense	31081	31081 BACE:1510L21 siRNA	UGGUUGGUAACCUCACCCATT	244
BACE	1753	UCACCUUGGACAUGGAAGACUGU	18	antisense	31082	31082 BACE:1773L21 siRNA	AGUCUUCCAUGUCCAAGGUTT	245
BACE	2457	CCUAACAUUGGUGCAAAGAUUGC	20	antisense	31083	31083 BACE:2477L21 siRNA	AAUCUUUGCACCAAUGUUATT	246
BACE	3583	UAUGGGACCUGCUAAGUGUGGAA	19	antisense	31084	31084 BACE:3603L21 siRNA	CCACACUUAGCAGGUCCCATT	247
BACE	2457	CCUAACAUUGGUGCAAAGAUUGC	8	sense	31378 E	31378 BACE:2459U21 sIRNA stab04	B uAAcAuuGGuGcAAAGAuuTT B	248
BACE	2457	CCUAACAUUGGUGCAAAGAUUGC	82	antisense	31381 E	31381 BACE:2477L21 siRNA	AAucuuuGcAccAAuGuuATsT	249
BACE	2457	CCUAACAUUGGUGCAAAGAUUGC	20	seuse	31384 E	31384 BACE:2459U21 siRNA	B uAAcAuuGGuGcAAAGAuuTT B	250
BACE	2457	CCUAACAUUGGUGCAAAGAUUGC	8	antisense	31387 E	BACE:2477L21 siRNA	AAucuuuGcAccAAuGuuATsT	251
BACE	2457	CCUAACAUUGGUGCAAAGAUUGC	20	seuse	31390 E	BACE:2459U21 siRNA inv	B uuAGAAAcGuGGuuAcAAuTT B	252
BACE	2457	CCUAACAUUGGUGCAAAGAUUGC	8	antisense	31393 B	antisense 31393 BACE:2477L21 siRNA	AuuGuAAccAcGuuucuAATsT	253
			1		4	Z459C) INV STADUS		

254	255	256	257	258	259	260	261	262	263	264	265	266	267	268	269	270	271	272	273	274	275	276
B uuAGAAAcGuGGuuAcAAuTT B	AuuGuAAccAcGuuucuAATsT	B GcuGucucuGAAGAcucuGTT B	B uuAcGuGGccuGuuucAAcTT B	B uuuGGAucAGGGAGuuGGATT B	cAGAGucuucAGAGAcAGcTsT	GuuGAAAcAGGccAcGuAATsT	uccAAcucccuGAuccAAATsT	GCUGUCUCUGAAGACUCUGTT	GGGAUGAUCAACAGGGUAGTT	UNACGUGGCCUGUUUCAACTT	UUUGGAUCAGGGAGUUGGATT	CAGAGUCUUCAGAGACAGCTT	CUACCCUGUUGAUCAUCCCTT	GUUGAAACAGGCCACGUAATT	UCCAACUCCCUGAUCCAAATT	B GGGAUGAUCAACAGGGUAGTT B	cuAccouGuuGAucAucccTsT	B GAUGGGACAACUAGUAGGGTT B	cccuAcuAGuuGucccAucTsT	B GGGAUGAUCAACAGGGUAGTT B	cuAcccuGuuGAucAuccTsT	B GAUGGGACAACUAGUAGGGTT B
31396 BACE:2459U21 siRNA inv stab07	31399 BACE:2477L21 siRNA (2459C) inv stab11	30737 BCL2:2100U21 siRNA stab04	30739 BCL2:4428U21 siRNA stab04	30740 BCL2:6233U21 sIRNA stab04	11 BCL2:2118L21 siRNA (2100C) stab05	30743 BCL2:4446L21 siRNA (4428C) stab05		30997 BCL2:2100U21 siRNA	30998 BCL2:3222U21 siRNA	30999 BCL2:4428U21 siRNA	31000 BCL2:6233U21 siRNA	31073 BCL2:2118L21 siRNA	31074 BCL2:3240L21 siRNA	antisense 31075 BCL2:446L21 siRNA (4428C)	31076 BCL2:6251L21 siRNA (6233C)	31368 BCL2:3222U21 siRNA stab04	31369 BCL2:3240L21 siRNA (3222C) stab05	31370 BCL2:3222U21 siRNA inv stab04	1 BCL2:3240L21 siRNA (3222C) inv stab05	31372 BCL2:3222U21 siRNA stab07	31373 BCL2:3240L21 siRNA (3222C) stab11	31374 BCL2:3222U21 siRNA inv stab07
3139		3073	3073	3074	30741		30744	3099	3099	3099	3100	3107	3107	3107		3136		3137	31371	3137		3137
seuse	antisense	seuse	seuse	seuse	antisense	antisense	antisense	sense	seuse	seuse	seuse	antisense	antisense	antisense	antisense	seuse	antisense	seuse	antisense	sense	antisense	seuse
20	70	21	22	23	21	22	23	21	24	22	23	12	24	22	23	24	24	24	24	24	24	24
CCUAACAUUGGUG	CCUAACAUUGGUGCAAAGAUUGC	UGGCUGUCUGAAGACUCUGCU	CUUUACGUGGCCUGUUUCAACAC				AGUUUGGAUCAGGGAGUUGGAAG		CAGGGAUGAUCAAC			ueecueucucueaagacucuecu	CAGGGAUGAUCAACAGGGUAGUG	CUUUACGUGGCCUGUUUCAACAC	AGUUUGGAUCAGGGAGUUGGAAG	CAGGGAUGAUCAACAGGGUAGUG	CAGGGAUGAUCAACAGGGUAGUG	CAGGGAUGAUCAACAGGGUAGUG	CAGGGAUGAUCAACAGGGUAGUG	CAGGGAUGAUCAACAGGGUAGUG	CAGGGAUGAUCAACAGGGUAGUG	CAGGGAUGAUCAACAGGGUAGUG
	2457	2098	4426	6231	2098	4426	6231	2098	3220	4426	6231	2098	3220	4426	6231	3220	3220	3220	3220	3220	3220	3220
BACE	BACE	BCL2	BCL2	BCL2	BCL2	BCL2	BCL2	BCL2	BCL2	BCL2	BCL2	BCL2	BCL2	BCL2	BCL2	BCL2	BCL2	BCL2	BCL2	BCL2	BCL2	BCL2

277	3 278	279	280	284		282	283	284	285	286	287	288	289	290	291	292		292	292	293	293	294	295	
cccuAcuAGuuGucccAucTsT	B uGuAGuGGGGuucuAGGcATT B	B AcAcAAccuucuGccuuuTT B	B AcAuuGuuuGcuGcuAuuGTT B	uGccuAGAAcccAcuAcATsT		AAAGGcAGAAGGuuuGuGuTsT	CAAUAGCAGCAAACAAUGUTsT	ACACUUCCUCUCCAAAAUGTT	UGUAGUGGGGUUCUAGGCATT	ACACAAACCUUCUGCCUUUTT	ACAUUGUUUGCUGCUAUUGTT	CAUUUUGGAGAGGAAGUGUTT	UGCCUAGAACCCCACUACATT	AAAGGCAGAAGGUUUGUGUTT	CAAUAGCAGCAAACAAUGUTT	B AcAcuuccucuccAAAAuGTT B		B AcAcunccucuccAAAAuGTT B	B AcAcuuccucuccAAAAuGTT B	cAuuuuGGAGAGGGAAGuGuTsT	cAuuuuGGAGAGGAAGuGuTsT	B GuAAAAccucuccuucAcATT B	uGuGAAGGAGGuuuuAcTsT	
antisense 31375 BCL2:3240L21 siRNA	30746 CCND1:1628U21 siRNA	30747 CCND1:2617U21 siRNA	30748 CCND1:3124U21 siRNA	30750 CCND1:1646L21 siRNA	(1628C) stab05	30751 CCND1:2635L21 siRNA (2617C) stab05	30752 CCND1:3142L21 siRNA (3124C) stab05	31009 CCND1:695U21 siRNA	31010 CCND1:1628U21 siRNA	31011 CCND1:2617U21 siRNA	31012 CCND1:3124U21 siRNA	31085 CCND1:713L21 siRNA (695C)	31086 CCND1:1646L21 siRNA	31087 CCND1:2635L21 siRNA	31088 CCND1:3142L21 siRNA	31304 CCND1:695U21 sIRNA	stab04	31304 CCND1:695U21 siRNA stab04	31304 CCND1:695U21 sIRNA stab04	31305 CCND1:713L21 siRNA (695C) stab05	31305 CCND1:713L21 siRNA	31316 CCND1:695U21 siRNA	31317 CCND1:713L21 siRNA	(695C) inv stab05
antisense	sense	sense	sense 3	antisense 3		antisense 3	antisense 3	sense 3	sense 3	sense 3	sense 3	antisense 3	antisense 3	antisense 3	antisense 3	sense 3	T	sense 3	sense 3.	antisense 3	antisense 3	sense 3.	antisense 31	
24	52	26	27	25	-	56	27	28	22	26	27	78	52	26	27	28	8	28	28	28	28	28	28	8
3220 CAGGGAUGAUCAACAGGGUAGUG	GCUGUAGUGGGGUUCUAGGCAUC	ACACACAAACCUUCUGCCUUUGA	UCACAUUGUUUGCUGCUAUUGGA	GCUGUAGUGGGGUUCUAGGCAUC	CONTRACTOR	_	UCACAUUGUUUGCU		GCUGUAGUGGGGUU		_1	_	GCUGUAGUGGGGUUCUAGGCAUC	ACACACAACCUUCUGCCUUUGA	UCACAUUGUUUGCUGCUAUUGGA	GAACACUUCCUCCAAAAUGCC		GAACACOOCCOCCAAAAOGCC	GAACACUUCCUCUCCAAAAUGCC	GAACACUUCCUCUCCAAAAUGCC	GAACACUUCCUCUCCAAAAUGCC	GAACACUUCCUCUCCAAAAUGCC	GAACACUUCCUCUCCAAAAUGCC	CHEGACACH GAGACH CACCACT
	1628	1 2617	3124	1646	2626	-+				_	1		1646	2635	3142	695	202		692	713	713	969	713	344
BCL2	CCND1	CCND1	CCND1	CCND1	CNO.		CCND	CCND	COND			ב ב	CCND1	CCND1	CCND1	CCND1	ניועט		CCND1	CCND1	CCND1	CCND1	CCND1	CDK2

297	298	299	300	301	302	303	304	305	306	307	308	309	310	311	312	313	314	315	316	317	318	319	320
AUCAAGCUAGCAGACUUUGTT	CUCACCUUCUAGUCUUGGCTT	ACGUUAGAUUUGCCGUACCTT	ACCCUCAGUCUCAGUGUCCTT	CAAAGUCUGCUAGCUUGAUTT	GCCAAGACUAGAAGGUGAGTT	GGUACGGCAAAUCUAACGUTT	B uGGucAcAGGAGAGAGGCTT B	B AGAAGuuGGGcuAucAAuGTT B	B uucAGGGGAcAuGAGuuuuTT B	GccuucucucaGuGAccATsT	cAuuGAuAGcccAAcuucuTsT	AAAAcucAuGucccuGAATsT	UGGUCACAGGAGAGGCTT	AGAAGUUGGGCUAUCAAUGTT	AGGGUGAUGGAUUGGAGUUTT	UUCAGGGGACAUGAGUUUUTT	GCCUUCUCUCUGUGACCATT	CAUUGAUAGCCCAACUUCUTT	AAGUCGAAUCCAUCACCCUTT	AAAACUCAUGUCCCCUGAATT	B AGGGuGAuGGAuuGGAGuuTT B	AAcuccAAuccAucAcccuTsT	B uuGAGGuuAGGuAGuGGGATT B
31566 CDK2:654U21 siRNA	31567 CDK2:1245U21 siRNA	31568 CDK2:1428U21 siRNA	31569 CDK2:362L21 siRNA (344C)	CDK2:672L21 siRNA (654C)	CDK2:1263L21 sIRNA (1245C)	31572 CDK2:1446L21 siRNA (1428C)	CHEK1:371U21 siRNA stab04	CHEK1:1351U21 siRNA stab04	CHEK1:1880U21 siRNA stab04	CHEK1:389L21 siRNA (371C) stab05	CHEK1:1369L21 siRNA (1351C) stab05	CHEK1:1898L21 siRNA (1880C) stab05	CHEK1:371U21 siRNA	CHEK1:1351U21 sIRNA	CHEK1:1492U21 siRNA	CHEK1:1880U21 siRNA	CHEK1:389L21 sIRNA (371C)	31078 CHEK1:1369L21 siRNA (1351C)		CHEK1:1898L21 siRNA (1880C)	CHEK1:1492U21 siRNA stab04	31303 CHEK1:1510L21 siRNA (1492C) stab05	CHEK1:1492U21 siRNA inv stab04
31566	31567	31568	31569	31570	31571	31572	30753	30754	30756	30757	30758		31001	31002	31003	31004	31077		31079	31080	31302		31314
sense	sense	sense	antisense	antisense	antisense	antisense	seuse	seuse	seuse	antisense	antisense	antisense	seuse	seuse	sense	sense	antisense	antisense	antisense	antisense	seuse	antisense	sense
30	31	32	29	30	31	32	33	34	35	33	34	35	33	34	36	35	33	. 34	36	35	36	36	36
CCAUCAAGCUAGCAGACUUUGGA		ACACGUUAGAUUUGCCGUACCAA	CUGGACACUGAGGGUGU	ccaucaagcuagcagacuuugga	CACUCACCUUCUAGUCUUGGCCA	ACACGUDAGAUUUGCCGUACCAA	UAUGGUCACAGGAGAGGCAA	UGAGAAGUUGGGCUAUCAAUGGA	GUUUCAGGGGACAUGAGUUUUCC	UAUGGUCACAGGAGAGAGGCAA	UGAGAAGUUGGGCUAUCAAUGGA	GUUUCAGGGGACAUGAGUUUUCC			UAAGGGUGAUGGAUUGGAGUUCA	<u> </u>	UAUGGUCACAGGAGAGAGGCAA	UGAGAAGUUGGGCUAUCAAUGGA	UAAGGGUGAUGGAUUGGAGUUCA	GUUUCAGGGGACAUGAGUUUUCC	UAAGGGUGAUGGAUUGGAGUUCA	UAAGGGUGAUGGAUUGGAGUUCA	UAAGGGUGAUGGAUUGGAGUUCA
654	1245	1428	362	672	1263	1446	369	1349	1878	369	1349	1878	369	1349	1490	1878	369	1349	1490	1878	1490	1490	1490
CDK2	CDK2	CDK2	CDK2	CDK2	CDK2	CDK2	СНЕК1	CHEK1	CHEK1	CHEK1	CHEK1	CHEK1	CHEK1	CHEK1	CHEK1	CHEK1	CHEK1	CHEK1	CHEK1	CHEK1	CHEK1	CHEK1	CHEK1

321	322	323	324	325	326	327	328	329	330	331	332	333	334	335	336
uccacuAccuAaccucAATsT	B UAACCUCGUACUGGUGCCUCC B	B GGAGGCACCAGUACGAGGUUA B	B AAACUCCAAGAUCCCCAAUCA B	B UGAUUGGGGAUCUUGGAGUUU B	B GCAAAACCCUGUGAUUUCCU B	B AGGAAAUCACAGGGUUUUUGC B	B UUGGUCAGUUCUGGCAGUUC B	B GAACUGCCAGAAACUGACCAA B	B CCUCCEUGGUCAUGCUCCAAU B	B AUUGGAGCAUGACCACGGAGG B	UAACCUCGUACUGGUGCCUCCUU	GGAGGCACCAGUACGAGGUUAUU	AAACUCCAAGAUCCCAAUCAUU	UGAUUGGGGAUCUUGGAGUUUUU	GCAAAAACCCUGUGAUUUCCUUU
antisense 31315 CHEK1:1510L21 sIRNA (1492C) inv stab05		antisense 25228 RPI 21550 EGFR 3830L23 AS as siRNA Str 2 (antisense)	25229 RPI 21549 EGFR as siRNA Str 2 (antisense)	25230 RPI 21549 EGFR 3 as siRNA Str 1 (sense)	25233 RPI 21545 EGFR as siRNA Str 2 (antisense)	25234 RPI 21545 EGFR as siRNA Str 1 (sense)	25235 RPI 21543 EGFR as siRNA Str 2 (antisense)	25236 RPI 21543 EGFR as siRNA Str 1 (sense)	25249 RPI 21550 EGFR 3830L23 AS as siRNA Str 1 (sense) Inverted Control	25250 RPI 21550 EGFR 3830L23 AS as siRNA Str 1 (sense) Inverted Control Compliment	25804 RPI 21550 EGFR 3830L23 AS as siRNA Str 1 (sense) +2U overhand	antisense 25805 RPI 21550 EGFR 3830L23 AS as sIRNA Str 2 (antisense) +2U overhand	antisense 25806 RPI 21549 EGFR as siRNA Str 2 (antisense)+	25807 RPI 21549 EGFR 3 as sIRNA Str 1 (sense)+2U overhand	antisense 25810 RPI 21545 EGFR as siRNA Str 2
3131	25227	2522		2523(25234	25235	25236	25249	25250	25804	25805	25806	25807	25810
antisense	seuse	antisense	antisense	sense	antisense	esues	antisense	sense	seuse	sense	seuse	antisense	antisense	sense	antisense
36	37	88	33	40	41	42	43	44	38	45	37	88	96 96	40	14
UAAGGGUGAUGGAL		ACCUCGUACUGGUGCCUCC	AUUGGGGAUCUUGGAGUUU	UGAUUGGGGAUCUUGGAGU	GAAAUCACAGGGUUUUUGC	AGGAAAUCACAGGGUUUUU	ACUGCCAGAACUGACCAA	GAACUGCCAGAAACUGACC	ACCUCGUACUGGUGCCUCC	AGGCACCAGUACGAGGUUA	UAACCUCGUACUGGUGCCU	ACCUCGUACUGGUGCCUCC	AUUGGGGAUCUUGGAGUUU	UGAUUGGGGAUCUUGGAGU	GAAAUCACAGGGUUUUUGC
	3828								3828	3828	3828				
CHEK1	EGFR	EGFR	EGFR	EGFR	EGFR	EGFR	EGFR			EGFR		EGFR	EGFR	EGFR	EGFR

s hang s +2U A Str ang	as as as e)+2U RNA Str srhang RNA Str srhang as as ense)+	rang Pang A Str A Str A Str (e)+	s hang s to the september of the septemb	A Str A Str ang	s s t-2U A Str ang se)+ se)+ E 2U E 2U E 2U E 2U E 2U E 2U	s s s s s s s s s s s s s s s s s s s	s hang hang se)+ se)+ se)+ se)+ se)+ se)+ se)+ se)+	s s t-2U A Str ang ang E 2U	S S S S S S S S S S S S S S S S S S S	S S S S S S S S S S S S S S S S S S S
Section Continued	25812 RPI 21543 EGFR as siRNA Str 2 (antisense)+2U overhang 25824 RPI 21550 EGFR as siRNA Str 1 (sense)+2U overhang 25825 RPI 21550 EGFR 3830L23 AS as siRNA 2 (antisense) +2U overhang	25 RPI 21550 EGFR as siRNA Str 2 (antisense)+20 overhang 1243 EGFR as siRNA Str 1 (sense)+20 overhang 1250 EGFR 3830L23 AS as siRNA Str 1 (sense) +20 overhang 25 RPI 21550 EGFR 3830L23 AS as siRNA Str 2 (antisense) +20 overhang 210 overhang 210 overhang 220 overhang 21 overhang 21 siRNA Str 1 (sense)+20 overhang 21 siRNA Str 1 (sense)+20 overhang 21 overhang 22 overhang 22 overhang 23 siRNA Str 1 (sense)+2 overhang 24 siRNA Str 1 (sense)+2 overhang 25 overhang 26 overhang 27 siRNA Str 1 (sense)+2 overhang overhang 27 siRNA Str 1 (sense)+2 overhang overhang 25 over	overhang RPI 21543 EGFR as siRNA Str 2 (antisense)+2U overhang RPI 21543 EGFR as siRNA Str 1 (sense)+2 overhang RPI 21550 EGFR 3830L23 AS as siRNA 1 (sense) +2U overhang RPI 21550 EGFR 3830L23 AS as siRNA 2 (antisense) +2U overhang RPI 21549 EGFR as siRNA Str 2 (antisense siRNA Str 2 (antisense siRNA Str 2 (sense)+2 overhang RPI 21549 EGFR as siRNA Str 1 (sense)+2 overhang RPI 21549 EGFR as siRNA Str 1 (sense)+2 overhang RPI 21549 EGFR as siRNA Str 1 (sense)+2 overhang RRI 21545 EGFR as siRNA Str 1 (sense)+2 overhang RRI 21545 EGFR as siRNA Str 1 (sense)+2 overhang	RPI 21543 EGFR as siRNA Str 2 (antisense)+20 overhang RPI 21543 EGFR as siRNA Str 1 (sense)+20 overhang Str 21550 EGFR as siRNA Str 1 (sense) +20 overhang siRNA Str 2 (antisense) +20 overhang Str 1 (sense)+20 overhang Str 2 (antisense)+20 overhang Str 1 (sense)+20 overhang Str 2 overhang	enhang 121543 EGFR as RNA Str 2 Intisense)+2U overhang 121550 EGFR Sense) +2U overhang 121550 EGFR 30L23 AS as siRNA 30L23 AS as siRNA 30L23 AS as siRNA 30L23 AS as siRNA 3121550 EGFR 30L23 AS as siRNA 3121550 EGFR 3121545 EGFR 32140 EGFR 3214	1243 EGFR as 12543 EGFR as 12543 EGFR as 12543 EGFR as 12543 EGFR as 12550 EGFR as 12549 EGFR as 12545 EGFR as 12555 EGFR as 125	12.15.15.15.15.15.15.15.15.15.15.15.15.15.	21543 EGFR as 1. (Sense) 7. (Sense) 7. (Sense) 7. (Sense) 7. (Sense) 4. (Sens	121543 EGFR as INA Str 2 121543 EGFR as INA Str 2 121543 EGFR as INA Str 2 121550 EGFR 121549 EGFR 121543 EGFR 121545 EGFR 121	INTERNATION (Serice) TO Serice) TO Serice) TO Serice) TO SERICE AS INDICATED (SERICE) SERICE AS INDICATED (SERICE) SERICE AS S
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الما ما ما	1 1 10 10 1	258 258 258 258	25813 25826 25826 25827 25827	25813 25824 25825 25826 25827 25830 2	Control Cont	Canton C	(anti 25833 RPI 25824 RPI 3833 3834 3834 25825 RPI 20 over 25830 RPI 31RN 25830 RPI 31RN 31RN 31RN 31RN 31RN 31RN 31RN 31RN	(anti) (anti) (b) 25821 RPI 2830 (c) 25825 RPI 2830 (c) 25825 RPI 2930 (c) 25830 RPI 2930 (c) 25831 RPI 2931 (c) 25831 RPI 2931 (c) 25831 RPI 2931 (c) 25831 RPI 2931 (c) 25833 RRIV (c) 2	(an 25813 RP 25824 RP 383 383 383 383 383 383 383 383 383 383 383 38705 SEB	25813 R 25825 3 3 3 3 3 3 3 3 3
									sense antisense antisense antisense sense sense sense sense sense sense sense	sense antisense antisense antisense sense
38 38	38 38	38 39 40	39 39 40 41	39 39 41 41	39 39 41 41 41 41 41 41	38 39 40 41 43 43	38 39 41 42 43 43 44	38 39 40 41 44 44 44	38 38 38 44 44 44 44 44 44 44 44 44 44 44 44 44	38 38 38 38 44 44 44 44 44 44 44 44 44 44 44 44 44
AUUGGGGAUCUUGGAGUUU	AUUGGGGAUCUUGGAGUUU UGAUUGGGGAUCURGGAGU	AUUGGGGAUCUUGGAGUUU UGAUUGGGGAUCUUGGAGU	AUUGGGGAUCUUGGAGUUU UGAUUGGGGAUCUUGGAGU GAAAUCACAGGGUUUUUGC	AUUGGGGGAUCUUGGAGUUU UGAUUGGGGAUCUUGGAGU GAAAUCACAGGGUUUUUGC	AUUGGGGAUCUUGGAGUUU UGAUUGGGGAUCUUGGGAGU GAAAUCACAGGGUUUUU AGGAAAUCACAGGGUUUUU	AUUGGGGAUCUUGGAGUUU UGAUUGGGGAUCUUGGAGU GAAAUCACAGGGUUUUUU AGGAAAUCACAGGGUUUUUU AGGAAAUCACAGGAAACUGACC	AUUGGGGGAUCUUGGAGUUU UGAUUGGGGGAUCUUGGC GAAAUCACAGGGUUUUUU AGGAAAUCACAGGGUUUUUU AGGAAAUCACAGGGUUUUUU GAACUGCCAGAAACUGACC	AUUGGGGAUCUUGGAGUUU UGAUUGGGGAUCUUGGGGAAAUCACAGGGUUUUUU AGGAAAUCACAGGGUUUUUU AGGAAAUCACAGGGUUUUUU AGGAAAUCACAGGGUUUUUU GAACUGCCAGAAACUGACC GAACUGCCAGAAACUGACC	AUUGGGGGAUCUUGGAGUUU UGAUUGGGGAUCUUGGAGU GAAAUCACAGGGUUUUUU AGGAAAUCACAGAAACUGACCAA GAACUGCCAGAAACUGACC GAACUGCCAGAAACUGACC AGGAAAUCACAGGGUUUUU	AUUGGGGGAUCUUGGAGUUU UGAUUGGGGGAUCUUGGC GAAAUCACAGGGUUUUUU ACUGCCAGAAACUGACCAA GAACUGCCAGAAACUGACC GAACUGCCAGAAACUGACC GAACUGCCAGAAACUGACC AGGAAAUCACAGGGUUUUU GUUCCGUGAGUUGAUCAUC
	5	40	41	41 41 42	41 41 42 43	43 43 43	43 43 44	44 44 44 44	44 44 45 42 42 44 44 44 44 44 44 44 44 44 44 44	44 44 44 46 46

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372	373	374	375	376	377	378	379	380	381	382	383	384	385	386	387	388	389	390	391	392	393	394	395	396
GGAGAuGGuuGAGcAGcuuTsT	cuGuGGAAGGAGGuuGTsT	GUGAAUGGCUCAAGGAACUTT	GGAACUGUGCAAGAUGACCTT	AAGCUGCUCAACCAUCUCCTT	CAACCAUCUCCAUCCACAGTT	AGUUCCUUGAGCCAUUCACTT	GGUCAUCUUGCACAGUUCCTT	GGAGAUGGUUGAGCAGCUUTT	CUGUGGAAGGAGAUGGUUGTT	CAUGCGACUGAGACAGCUCTT	ACAUCCUGACUUCUGUGAGTT	GAUGAUGAUGAGAGACGTT	ACAAUUUCUGUGCCAUUGCTT	GAGCUGUCAGUCGCAUGTT	CUCACAGAAGUCAGGAUGUTT	CGUCUCCAUCAUCAUCTT	GCAAUGGCACAGAAAUUGUTT	CsUsGsAsGsUUUAAAAGGCACCCTsT	CSASASCSCSACAAAAUACAACAATST	CsCsUsGsGsAAAGAAUCAAAACCTsT	GSCsAsAsGsGAGGGCCUCUGAUGTsT	GSGSGSUSGSCCUUUUAAACUCAGTST	Ususesususeuauuuugueeuugetst	GsGslislislislical III III III III III III III III III I
antisense 30767 ERG2:779L21 siRNA (761C) stab05		31045 ERG2:244U21 siRNA	31046 ERG2:519U21 siRNA	31047 ERG2:761U21 siRNA	31048 ERG2:769U21 siRNA	31121 ERG2:262L21 siRNA (244C)	31122 ERG2:537L21 siRNA (519C)	23 ERG2:779L21 siRNA (761C)	31124 ERG2:787L21 sIRNA (769C)	16 EZH2:203U21 siRNA	17 EZH2:340U21 siRNA	31418 EZH2:690U21 siRNA	31419 EZH2:1495U21 SIRNA	31420 EZH2:221L21 sIRNA (203C)	21 EZH2:358L21 siRNA (340C)	31422 EZH2:708L21 siRNA (690C)	23 EZH2:1513L21 siRNA (1495C)	29694 FLT1:349U21 siRNA stab01	29695 FLT1:2340U21 siRNA stab01	29696 FLT1:3912U21 siRNA stab01	97 FLT1:2949U21 siRNA stab01	98 FLT1:369L21 siRNA (349C) stab01		antisense 29700 El T1-39321 21 siRNIA
se 307	зе 30768	\vdash						31123		31416	31417		314		314	e 314	e 31423	2966	2966	2966	29697	e 29698	е 29699	2076
antisens	antisense	seuse	seuse	seuse	sense	antisense	antisense	antisense	antisense	sense	esues	seuse	esues	antisense	antisense 31421	antisense	antisense	esues	seuse	sense	seuse	antisense	antisense	antisens
20	51	84	49	20	51	48	49	20	51	52	23	24	22	52	23	54	55	56	25	28	29	29	27	58
GAAAGCUGCUCAACCAUCUCCUU	CUCAACCAUCUCCUUCCACAGUG	AGGUGAAUGGCUCAAGGAACUCU	AAGGAACUGUGCAAGAUGACCAA	GAAAGCUGCUCAACCAUCUCCUU	CUCAACCAUCUCCUUCCACAGUG	AGGUGAAUGGCUCAAGGAACUCU	AAGGAACUGUGCAAGAUGACCAA	GAAAGCUGCUCAACCAUCUCCUU	cucaaccaucuccuuccacaeue	UACAUGCGACUGAGACAGCUCAA				UACAUGCGACUGAGACAGCUCAA	GCACAUCCUGACUUCUGUGAGCU	ACGAUGAUGAUGGAGACGAU	ugacaauuucugugccauugcua	AACUGAGUUUAAAAGGCACCCAG		AGCCUGGAAAGAAUCAAAACCUU	AAGCAAGGGCCUCUGAUGGU	AACUGAGUUUAAAAGGCACCCAG	AACAACCACAAAAUACAACAAGA	AGCCUGGAAAGAAUCAAAACCUU
759	767	242	517	759	767	242	517	759	767	201	338	688	1493	201	338	889	1493	347	2338	3910	2947	347	2338	3910
ERG2	ERG2	ERG2	ERG2	ERG2	ERG2	ERG2	ERG2	ERG2	ERG2	EZH2	EZH2	EZH2	EZH2	ЕХН2	ЕZН2	EZH2	EZH2	FLT1	FLT1	FLT1	FLT1	FLT1	FLT1	FLT1

					(3912C) stab01	
AAGCAAGGAGGCCUCUGAUGGU	AAGGAGGCCUCUGAUGGU	29	antisense		29701 FLT1:2969L21 siRNA (2949C) stab01	CSASUSCSASGAGGCCCUCCUUGCTST
AACUGAGUUUAAAAGGCACCCAG	IGAGUUUAAAAGGCACCCAG	26	seuse	29702	29702 FLT1:349U21 siRNA stab03	csusGsAsGuuuAAAAGGcAcscscsTsT
AACAACCACAAAAUACAACAAGA		21	seuse	29703	29703 FLT1:2340U21 siRNA stab03	csAsAscsCAcAAAuAcAAcsAsAsTsT
CAAAACCUU	CAAAACCUU	28	seuse	29704	29704 FLT1:3912U21 siRNA stab03	cscsusGsGAAAGAAucAAAAscscsTsT
AAGCAAGGAGGCCUCUGAUGGU 59	ucugaugeu	I —	seuse	29705	29705 FLT1:2949U21 siRNA stab03	GscsAsAsGGAGGGccucuGAsusGsTsT
AACUGAGUUUAAAAGGCACCCAG 5	<u> </u>	20	antisense		29706 FLT1:369L21 siRNA (349C) stab02	GsGsGsUsGsCsCsUsUsUsUsAsAsAsCsUs CsAsGsTsT
AACAACCACAAAAUACAACAAGA 57	ACAACAAGA	I	antisense		29707 FLT1:2358L21 siRNA (2340C) stab02	UsUsGsUsUsGsUsAsUsUsUsGsUsGsG sUsUsGsTsT
AGCCUGGAAAGAAUCAAAACCUU 58	CAAAACCUU	l 1	antisense		29708 FLT1:3932L21 siRNA (3912C) stab02	GsGsUsUsUsGsAsUsUsCsUsUsUsCsCs AsGsGsTsT
AAGCAAGGAGGCCUCUGAUGGU 59	JCUGAUGGU	_	antisense		29709 FLT1:2969L21 siRNA (2949C) stab02	CsAsUsCsAsGsAsGsGsCsCsUsCsCsUs UsGsCsTsT
AACAACCACAAAAUACAACAAGA 57		١.	seuse	29981	29981 FLT1:2340U21 siRNA Native	CAACCACAAAUACAACAAGA
AACAACCACAAAAUACAACAAGA 57			antisense		29982 FLT1:2358L21 siRNA (2340C) Native	ບບຣບັບຣບAບບບບຣບຣຣບບຣບບ
AACAACCACAAAAUACAACAAGA 57			seuse	29983	29983 FLT1:2342U21 siRNA stab01 inv	Asascabaaaacaccaactst
AACAACCACAAAAUACAACAAGA 57	CAACAAGA	ī	antisense		29984 FLT1:2358L21 siRNA (2340C) stab01 inv	GSUSUSGSGSUGUUUNAUGUUGUUTST
AACAACCACAAAAUACAACAAGA 57			sense	29985	29985 FLT1:2342U21 siRNA stab03 inv	AsAscsAsAcAuAAAcAccAsAscsTsT
AACAACCACAAAAUACAACAAGA 57			antisense		29986 FLT1:2358L21 siRNA (2340C) stab02 inv	GsUsUsGsGsUsGsUsUsUsAsUsGsUsU sGsUsUsTsT
AACAACCACAAAAUACAACAAGA 57			seuse	29987	29987 FLT1:2340U21 siRNA inv Native	AGAACACAUAAAACACCAAC
AACAACCACAAAAUACAACAAGA 57			antisense		FLT1:2358L21 siRNA (2340C) inv Native	UNGUNGGNGNNNANGNNGNN
	H	١. ا	sense	30075	30075 FLT1:2340U21 siRNA	CAACCACAAAUACAACAATT
AACAACCACAAAAUACAACAAGA 57	ACAACAAGA	I	antisense	30076	antisense 30076 FLT1:2358L21 siRNA (2340C)	UUGUUGUAUUUUGUGGUUGTT
L	ACAACAAGA	7	sense	30077	30077 FLT1:2342U21 siRNA Inv	AGAACACAUAAAACACCATT
AACAACCACAAAAUACAACAAGA 57	ACAACAAGA	_	antisense		30078 FLT1:2358L21 siRNA (2340C) inv	UVGUVGGUGUUUNAVGUVGTT
AACAACCACAAAUACAACAAGA 57	ACAACAAGA	~	antisense	30187	antisense 30187 FLT1:2358L21 siRNA	uuGuuGuAuuuuGuGGuuGTT

			L			10340C) 2'E 11 C		
FLT1	2338	AACAACCACAAAAUACAACAAGA	57	anticence	30100	aptisense 30100 El 71:03681 01 cibbin		
F	-+		5	a constant	00100	(2340C) nitroindole	nnennenvendendexX	419
=	7338	AACAACCACAAAAUACAACAAGA	22	antisense	30193		uuGuuGuAuuuuGuGGuuGZZ	420
FLT1	2338	AACAACCACAAAAUACAACAAGA	57	sense	30196	30196 FLT1:2340U21 siRNA	B CAACCACAAAAIACAACTT B	101
<u> </u>	2338					sense iB caps w/2'FY's		7
	-		27	sense	30199	FLT1:2340U21 siRNA sense iB caps	CAACCACAAAUACAACAATT	422
FLT			25	antisense	30340	30340 FLT1:2358L21 siRNA	uuGuuGuAuuuuGuGGuuGTX	423
FLT	2338	AACAACCACAAAAUACAACAAGA	23	antisense	30341	FLT1:2358L21 siRNA	uuGuuGuAuuuuGuGGuuGTX	424
FLT1	2338	AACAACCACAAAAUACAACAAGA	22	antisense		30342 FLT1:2358L21 siRNA	uuGuuGuAuuuuGuGGuuGTU	425
FLTI	2338	AACAACCACAAAAUACAACAAGA	22	antisense	30343	30343 FLT1:2358L21 siRNA	uuGuuGuAuuuuGuGGuuGTt	426
FLT1	2338	AACAACCACAAAAUACAACAAGA	22	antisense		30344 FLT1:2358L21 siRNA	uuGuuGuAuuuuGuGGuuGTu	427
FLT1	2338	AACAACCACAAAAUACAACAAGA	22	antisense	30345	30345 FLT1:2358L21 siRNA	uuGuuGuAuuuuGuGGuuGTD	428
FLT1	2338	AACAACCACAAAAUACAACAAGA	25	antisense	30346	30346 FLT1:2358L21 siRNA	uuGuuGuAuuuuGuGGuuGXT	429
FLT1	2338	AACAACCACAAAAUACAACAAGA	22	antisense	30416	FLT1:2358L21 siRNA	uuGuuGuAuuuuGuGGuuGTsT	430
FLT1	1182	1182 UCGUGUAAGGAGUGGACCAUCAU	90	sense	30777	FLT1:1184U21 siRNA	B GuGuAAGGAGGAccAucTT B	431
FLT1	3501	3501 UUACGGAGUAUUGCUGUGGGAAA	61	sense	30778 F	FLT1:3503U21 siRNA	B AcGGAGuAuuGcuGuGGGGATT B	432
FLT1	4713	UAGCAGGCCUAAGACAUGUGAGG	62	seuse	30779 F	30779 FLT1:4715U21 sIRNA	B GCAGGccuAAGACAuGuGATT B	433
FLT1	4751	AGCAAAAAGCAAGGGAAAAGA	63	seuse	30780 F	30780 FLT1:4753U21 siRNA	B cAAAAAGCAAGGGAGAAAATT B	434
FLT1	1182	1182 UCGUGUAAGGAGUGGACCAUCAU	8	antisense	30781 F	FLT1:1202L21 siRNA	GAuGGuccAcuccuuAcAcTsT	435
FLT1	3501	UNACGGAGUAUUGCU	61	antisense	30782 F	30782 FLT1:3521L21 siRNA (3503C) stab05	ucccAcAGcAAuAcuccGuTsT	436
FLT1	4713		62	antisense	30783 F	FLT1:4733L21 siRNA	ucAcAuGucuuAGGccuGcTsT	437
	4751		63	antisense	30784 F	30784 FLT1:4771L21 siRNA (4753C) stab05	unnucucconuGcunuuuGTsT	438
	2338	AACAACCACAAAAUACAACAAGA	22	seuse	30955 F	30955 FLT1:2340U21 siRNA	B CAACCACAAAUACAACAATT B	439

	440	441	442	443	444	445	446	447	448	449	450	451	452	453	454	455	456	457	458	459	460	461
	uuGuuGuAuuuuGuGGuuGTsT	AACAACAUAAAACACCAACTT	GUUGGUGUUUNAUGUUGUUTT	B AACAACAUAAAACACCAACTT B	GuuGGuGuuunAuGuuGuuTsT	B AACAACAUAAAACACCAACTT B	GuuGGuGuuudAuGuuGuuTsT	CUGAGUUUAAAAGGCACCCTT	GCAAGGAGGCCUCUGAUGTT	CCUGGAAAGAAUCAAAACCTT	GGGUGCCUUUNAAACUCAGTT	CAUCAGAGGCCCUCCUUGCTT	GGUUUUGAUUCUUUCCAGGTT	B cuGAGuuuAAAAGGcAccTT B	B GcAAGGAGGccucuGAuGTT B	B ccuGGAAAGAAucAAAAccTT B	GGGuGccuuuuAAAcucAGTsT	cAucAGAGGcccuccuuGcTsT	GGuuuuGAuucuuuccAGGTsT	B cuGAGuuuAAAAGGcAccCTT B	B GcAAGGAGGccucuGAuGTT B	B ccuGGAAAGAAucAAAAccTT B
stab07	30956 FLT1:2358L21 siRNA (2340C) stab08		30964 FLT1:2358L21 siRNA (2340C) inv	30965 FLT1:2340U21 siRNA stab04 inv	antisense 30966 FLT1:2358L21 siRNA (2340C) stab05 inv		30968 FLT1:2358L21 siRNA (2340C) stab08 inv	31182 FLT1:349U21 siRNA TT	31183 FLT1:2949U21 siRNA TT	31184 FLT1:3912U21 siRNA TT	31185 FLT1:367L21 siRNA (349C) TT	31186 FLT1:2967L21 siRNA (2949C) TT		31188 FLT1:349U21 siRNA stab04	31189 FLT1:2949U21 siRNA stab04	31190 FLT1:3912U21 siRNA stab04	1 FLT1:367L21 siRNA (349C) stab05				31195 FLT1:2949U21 siRNA stab07	31196 FLT1:3912U21 siRNA
		30963		3096	3096	30967		3118	3118	3118			31187	3118	3118	3119	3119	31192	31193	31194	31199	31196
	antisense	seuse	antisense	seuse	antisense	seuse	antisense	seuse	seuse	sense	antisense	antisense	antisense	seuse	sense	sense	antisense 31191	antisense	antisense	seuse	seuse	sense
	22	22	22	57	22	22	22	26	23	28	26	29	28	56	59	58	29	59	28	26	29	28
	AACAACCACAAAAUACAACAAGA	AACAACCACAAAAUACAACAAGA			AACAACCACAAAAUACAACAAGA	AACAACCACAAAAUACAACAAGA	AACAACCACAAAAU	AACUGAGUUUAAAAGGCACCCAG						AACUGAGUUUAAAAGGCACCCAG	_		AACUGAGUUUAAAAGGCACCCAG	AAGCAAGGAGGGCCI		AACUGAGUUUAAAAGGCACCCAG	-	AGCCUGGAAAGAAUCAAAACCUU
_	2338	2338	2338	2338	2338	2338	2338	347	2947	3910	347	2947	3910	347	2947	3910	347	2947	3910	347	2947	3910
	FLT1	FLT1	FLT1	FLT1	FLT1	FLT1	FLT1	FLT1	FLT1	FLT1	FLT1	FLT	FLT1	FLT1	FLT1	FLT1	FLT1	FLT1	FLT1	FLT1	FLT1	FLT1

462	463	464	465	466	467	468	469	470	471	472	473	474	475	476	477	478	479	480	481	482
GGGuGccuuuuAAAcucAGTsT	cAucAGAGGcccuccuuGcTsT	GGuuuuGAuucuuuccAGGTsT	CCCACGGAAAAUUUGAGUCTT	GUAGUCCCGGGAGGAACGTT	CCAAAACUAAGAAAGGUCCTT	GACUCAAAUUUUCCGUGGGTT	CGUUCCUCCGGAGACUACTT	GGACCUUUCUUAGUUUUGGTT	B cccAcGGAAAAuuuGAGucTT B	B GuAGucuccGGGAGGAAcGTT B	B ccAAAAcuAAGAAAGGuccTT B	GAcucAAAuuuuccGuGGGTsT	cGuuccuccGGAGAcuAcTsT	GGAccuuucuuAGuuuuGGTsT	B cccAcGGAAAuuuGAGucTT B	B GuAGucuccGGGAGGAAcGTT B	B ccAAAAcuAAGAAAGGuccTT B	GAcucAAAuuuuccGuGGGTsT	cGuuccuccGGAGAcuAcTsT	GGAccuuucuuAGuuuuGGTsT
antisense 31197 FLT1:367L21 siRNA (349C) stab08	31198 FLT1:2967L21 siRNA (2949C) stab08	31199 FLT1:3930L21 siRNA (3912C) stab08	31200 FLT1:349U21 siRNA inv TT	31201 FLT1:2949U21 siRNA inv TT	31202 FLT1:3912U21 siRNA inv	31203 FLT1:367L21 sIRNA (349C) inv TT	31204 FLT1:2967L21 siRNA (2949C) inv TT	antisense 31205 FLT1:3930L21 siRNA (3912C) inv TT	31206 FLT1:349U21 siRNA stab04 inv	31207 FLT1:2949U21 siRNA stab04 inv	31208 FLT1:3912U21 siRNA stab04 inv	31209 FLT1:367L21 siRNA (349C) stab05 inv	31210 FLT1:2967L21 siRNA (2949C) stab05 inv	31211 FLT1:3930L21 siRNA (3912C) stab05 inv	31212 FLT1:349U21 siRNA stab07 inv	31213 FLT1:2949U21 siRNA stab07 inv	31214 FLT1:3912U21 siRNA stab07 inv	31215 FLT1:367L21 siRNA (349C) stab08 inv	31216 FLT1:2967L21 siRNA (2949C) stab08 inv	
se 311						se 312	se 312	se 312												se 312
antisen	antisense	antisense	seuse	seuse	sense	antisense	antisense	antisen	seuse	seuse	seuse	antisense	antisense	antisense	seuse	seuse	seuse	antisense	antisense	antisense 31217
26	29	85	26	29	28	26	69	28	26	29	28	20	29	82	99	29	28	26	29	28
347 AACUGAGUUUAAAAGGCACCCAG	-	1 1	AACUGAGUUUAAAAGGCACCCAG	2947 AAGCAAGGAGGGCCUCUGAUGGU	AGCCUGGAAAGAAUCAAAACCUU				AACUGAGUUDAAAAGGCACCCAG	•		AACUGAGUUUAAAAGGCACCCAG		AGCCUGGAAAGAAUCAAAACCUU	AACUGAGUUNAAAAGGCACCCAG	2947 AAGCAAGGAGGGCCUCUGAUGGU		AACUGAGUUUAAAAGGCACCCAG		AGCCUGGAAAGAAUCAAAACCUU
347	2947	3910	347	2947	3910	347	2947	3910	347	2947	3910	347		3910	347	2947	3910	347	2947	3910
FLT1	FLT1	FLT1	FLT1	FLT1	FLT1	FLT1	FLT1	FLT1	FLT1	FLT1	FLT1	FLT1	FLT1	FLT1	FLT1	FLT1	FLT1	FLT1	FLT1	FLT1

483	484	485	486	487	488	489	490	491	492	493	494	495	496	497	498	499	200	501	502	503	504
B CUGAGUUUAAAAGGCACCCTT B	B GCAAGGAGGCCUCUGAUGTT B	B CCUGGAAAGAAUCAAAACCTT B	GGGUGCCUUUUAAACUCAGTST	CAUCAGAGGCCCUCCUUGCTsT	GGUUUUGAUUCUUUCCAGGTST	B CCCACGGAAAUUUGAGUCTT B	B GUAGUCUCCGGGAGGAACGTT B	B CCAAAACUAAGAAAGGUCCTT B	GACUCAAAUUUUCCGUGGGTsT	CGUUCCUCCGGAGACUACTST	GGACCUUUCUUAGUUUUGGTST	uuGuuGuAuuuuGuGGuuGXsX	cAucAGAGGccuccuuGcXsX	uuGuuGuAuuuuGuGGuuGXsT	cAucAGAGGcccuccuuGcXsT	B CAACCACAAAUACAACAATT B	B AACAACAUAAAACACCAACTT B	UUGUUGUAUUUUGUGGUUGTST	GUUGGUGUUUAUGUUGUUTST	B cAAcuGAGAAGccAAGAcuTT B	B cAuGGAccuAucuGGGuccTT B
	271 FLT1:2949U21 siRNA stab09	31272 FLT1:3912U21 siRNA stab09	273 FLT1:367L21 siRNA (349C) stab10	antisense 31274 FLT1:2967L21 siRNA (2949C) stab10	31275 FLT1:3930L21 sIRNA (3912C) stab10		31277 FLT1:2949U21 siRNA stab09 inv	78 FLT1:3912U21 siRNA stab09 inv	31279 FLT1:367L21 siRNA (349C) stab10 inv		31281 FLT1:3930L21 siRNA (3912C) stab10 inv	_	25 FLT1:2967L21 siRNA (2949C) stab11 3'-BrdU	42 FLT1:2358L21 siRNA (2340C) stab11 3'-BrdU	43 FLT1:2967L21 siRNA (2949C) stab11 3'-BrdU	31449 FLT1:2340U21 siRNA stab09	31450 FLT1:2340U21 siRNA inv stab09	31451 FLT1:2358L21 siRNA (2340C) stab10	31452 FLT1:2358L21 siRNA (2340C) inv stab10	30769 FOS:19U21 sIRNA stab04	30770 FOS:1028U21 siRNA
	31271	312	e 31273	912		31276	312	31278		e 31280		e 31424	e 31425	e 31442	e 31443	314	314			307	307
seuse	seuse	esues	antisense	antisens	antisense	seuse	seuse	seuse	antisense	antisense	antisense	antisense	antisense	antisense	antisense	esues	seuse	antisense	antisense	seuse	sense
56	29	28	29	29	28	26	29	28	26	29	28	22	29	25	29	22	22	22	22	64	65
	-			_	AGCCUGGAAAGAAUCAAAACCUU	AACUGAGUUUAAAAGGCACCCAG	-		AACUGAGUUUAAAAGGCACCCAG	AAGCAAGGAGGCCUCUGAUGGU			4		1		AACAACCACAAAAUACAACAAGA		_	AGCAACUGAGAAGCCAAGACUGA	
347	2947	3910	347	2947	3910	347	2947	3910	347	2947	3910	2338	2947	2338	2947	2338	2338	2338	2338	17	1026
FLT1	FLT1	FLT1	FLT1	FLT1	FLT1	FLT1	FLT1	FLT1	FLT1	FLT1	FLT1	FLT1	FLT1	FLT1	FLT1	FLT1	FLT1	FLT1	FLT1	FOS	FOS

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FOS 14	1403	UAGGGAGGACCUUAUCUGUGCGU	99	seuse	30771		B GGGAGGAccuuAucuGuGcTT B	505
FOS 1	1460	AAGCAUCCAUGUGUGGACUCAAG	29	seuse	30772		B GcAuccAuGuGuGGAcucATT B	506
FOS	17	AGCAACUGAGAAGCCAAGACUGA	64	antisense		30773 FOS:37L21 siRNA (19C) stab05	AGucuuGGcuucucAGuuGTsT	507
FOS 16	1026	GACAUGGACCUAUCUGGGUCCUU	65	antisense	30774		GGAcccAGAuAGGuccAuGTsT	508
FOS 14	1403	UAGGGAGGACCUUAUCUGUGCGU	99	antisense	30775		GcAcAGAuAAGGuccucccTsT	209
FOS 14	1460	AAGCAUCCAUGUGUGGACUCAAG	29	antisense	30776		uGAGuccAcAuGGAuGcTsT	510
FOS .	17	AGCAACUGAGAAGCCAAGACUGA	49	sense	31049	31049 FOS:19U21 siRNA	CAACUGAGAAGCCAAGACUTT	511
 	970	1026 GACAUGGACCUAUCUGGGUCCUU	65	sense	31050	31050 FOS:1028U21 siRNA	CAUGGACCUAUCUGGGUCCTT	512
\dashv		1403 UAGGGAGGACCUUAUCUGUGCGU	99	sense	31051	31051 FOS:1405U21 siRNA	GGGAGGACCUUAUCUGUGCTT	513
\dashv	_	AAGCAUCCAUGUGGACUCAAG	29	seuse	31052	31052 FOS:1462U21 siRNA	GCAUCCAUGUGGGACUCATT	514
FOS	1		64	antisense	31125	31125 FOS:37L21 siRNA (19C)	AGUCUUGGCUUCUCAGUUGTT	515
	1026	GACAUGGACCUAUCUGGGUCCUU	65	antisense	31126	31126 FOS:1046L21 siRNA (1028C)	GGACCCAGAUAGGUCCAUGTT	516
FOS 14	1403	UAGGGAGGACCUUAUCUGUGCGU	99	antisense		31127 FOS:1423L21 sIRNA (1405C)	GCACAGAUAAGGUCCUCCCTT	517
	1460	AAGCAUCCAUGUGGACUCAAG	29	antisense	31128	FOS:1480L21 sIRNA (1462C)	UGAGUCCACACAUGGAUGCTT	518
	2681	UGAAGAGGGAAAGCUGACAUCUG	89	sense	31541	31541 GAB2:2681U21 siRNA	AAGAGGGAAAGCUGACAUCTT	519
	4316	GAGGAAGGAAGGAGAGGCUU	69	sense	31542	31542 GAB2:4316U21 siRNA	GGAAGAAGGAAGGAGGCTT	520
	2006	GAGAGGACUGAGCCUACGGAAAG	20	esues	31543	31543 GAB2:5006U21 siRNA	GAGGACUGAGCCUACGGAATT	521
GAB2 59	5958	nunecneneeneveveveneenve	7.1	sense	31544	31544 GAB2:5958U21 siRNA	UGCUGUGGUGACACAUGGUTT	522
	2699	UGAAGAGGGAAAGCUGACAUCUG	89	antisense	31545	31545 GAB2:2699L21 siRNA (2681C)	GAUGUCAGCUUUCCCUCUUTT	523
GAB2 4		GAGGAAGGAAGGAGGCUU	69	antisense	31546	GAB2:4334L21 siRNA (4316C)	GCCUCUCCUUCCUUCCTT	524
_	5024	GAGAGGACUGAGGCCUACGGAAAG	02	antisense	31547	GAB2:5024L21 siRNA (5006C)	UUCCGUAGGCUCAGUCCUCTT	525
GAB2 59	5976	UUUGCUGUGGUGACACAUGGUAC	71	antisense	31548	GAB2:5976L21 siRNA (5958C)	ACCAUGUGUCACCACAGCATT	526
Her2		CCGCAGUGAGCACCAUGGA	72	antisense		25245 RPI 17763 Her2Neu AS as siRNA Str 2 (antisense)	B UCCAUGGUGCUCACUGCGGCU B	527
Her2		AGCCGCAGUGAGCACCAUG	73	seuse	25246	RPI 17763 Her2Neu AS as siRNA Str 1 (sense)	B AGCCGCAGUGAGCACCAUGGA B	528
Her2		CCGCAGUGAGCACCAUGGA	72	seuse	25247	25247 RPI 17763 Her2Neu AS	B AGGUACCACGAGUGACGCCGA B	529

					as siRNA Str 1 (sense)		
	caugeugeucacugegeeu	42	sense	25248	25248 RPI 17763 Her2Neu AS as siRNA Str 1 (sense) Inverted control	B UCGGCGUCACUCGUGGUACCU B	530
	CCGCAGUGAGCACCAUGGA	72	antisense	25822	antisense 25822 RPI 17763 Her2Neu AS as si 1776 Str 2 (antisense) 2U overhann	uccaugeucacueceecuuu	531
	AGCCGCAGUGAGCACCAUG	73	seuse	25823	25823 RPI 17763 Her2Neu AS as siRNA Str 1 (sense)+21 overhand	AGCCGCAGUGAGCACCAUGGAUU	532
	CCGCAGUGAGCACCAUGGA	2.	antisense	25842	antisense 25842 RPI 17763 Her2Neu AS as sIRNA Str 2 (antisense)+21 overhand	B UCCAUGGUGCUCACUGCGGCUUU B	533
	AGCCGCAGUGAGCACCAUG	73	seuse	25843	25843 RPI 17763 Her2Neu AS as siRNA Str 1 (sense)+2U overhand	B AGCCGCAGUGAGCACCAUGGAUU B	534
_	UGGGGUCGUCAAAGACGUU	75	sense	28262	28262 Her2.1.sense Str1	UGGGGUCGUCAAAGACGUUTT	535
3/06	UGGGGUCGUCAAAGACGUU	75	antisense	28263	28263 Her2.1.antisense Str2	AACGUCUUUGACGACCCCATT	536
	UGGGGUCGUCAAAGACGUU	75	sense	28264	28264 Her2.1.sense Str1 inverted	UUGCAGAACUGCUGGGGUTT	537
3706	UGGGGUCGUCAAAGACGUU	75	antisense		28265 Her2.1.antisense Str2 inverted	ACCCCAGCAGUUUCUGCAATT	538
	GGUGCUUGGAUCUGGCGCU	92	seuse	28266	28266 Her2.2.sense Str1	GGUGCUUGGAUCUGGCGCUTT	539
	GGUGCUUGGAUCUGGCGCU	76	antisense	28267	antisense 28267 Her2.2.antisense Str2	AGCGCCAGAUCCAAGCACCTT	540
	GeueconecAncocccco	76	sense	28268	28268 Her2.2.sense Str1 inverted	UCGCGGUCUAGGUUCGUGGTT	541
	GGUGCUUGGAUCUGGCGCU	76	antisense	28269	Her2.2.antisense Str2 inverted	CCACGAACCUAGACCGCGATT	542
	GAUCUUUGGGAGCCUGGCA	77	sense	28270	28270 Her2.3.sense Str1	GAUCUUUGGGAGCCUGGCATT	543
	GAUCUUUGGGAGCCUGGCA	2	antisense	28271	antisense 28271 Her2.3.antisense Str2	UGCCAGGCUCCCAAAGAUCTT	544
	GAUCUUUGGGAGCCUGGCA	4	sense	28272 + ii	28272 Her2.3.sense Str1 inverted	ACGGUCCGAGGGUUUCUAGTT	545
	GAUCUUUGGGAGCCUGGCA	22	antisense	28273 F	28273 Her2.3.antisense Str2 inverted	CUAGAAACCCUCGGACCGUTT	546
	GGUGCUUGGAUCUGGCGCU	9/	seuse	29989 F	29989 Her2.2.sense Str1 (site 2344)	GsGsusGscuuGGAucuGGcGscsusTsT	547
ြ	GGUGCUUGGAUCUGGCGCU	76	antisense	29990 F	antisense 29990 Her2.2.antisense Str2	AsGsCsGsCAGAUCCAAGCACCTsT	548
ا	GGUGCUUGGAUCUGGCGCU	92	sense	29991 H	29991 Her2.2.sense Str1 (site 2344)	GsGsUsGsCsUUGGAUCUGGCGCUTsT	549
							_

571	572	573	574	575	929	27.2	565	578	579	580	999	581	582	583	584	561	562	585	586	587	588	589
B GGuGcuuGGAucuGGcGcuTT B	B ucGcGGucuAGGuucGuGGTT B	B GGuGcuuGGAucuGGcGcuTT B	B AAcGucuuuGAcGAcccATT B	AGcGccAGAuccAAGcAccTsT	uGGGGucGucAAAGAcGuuTsT	B GAAuGGcucAGuGAccuGuTT B	B GGuGcuuGGAucuGGcGcuTT B	B AAcGucuuuGAcGAcccATT B	B cAccuucAAAGGGAcAccuTT B	AcAGGucAcuGAGccAuucTsT	AGcGccAGAuccAAGcAccTsT	uGGGGucGucAAAGAcGuuTsT	AGGuGucconuuGAAGGuGTsT	B uGGGGucGucAAAGAcGuuTT B	AAcGucuuuGAcGAcccATsT	B uGGGGucGucAAAGAcGuuTT B	AAcGucuuuGAcGAcccATsT	ACCAUUUUGUGGACGAAUATT	CUGUUGGACAUCCUGGAUATT	GGAUGCCUUCUACACGUUGTT	GAACCCUCCUGAUGAGAGUTT	UAUUCGUCCACAAAAUGGUTT
30448 Her2 sense (site 2344) stab6	30449 Her2 sense inverted (site 2344) stab6		30646 HER2:3726L21 siRNA		30648 HER2:3708U21 siRNA		30698 HER2:2346U21 siRNA stab04	30699 HER2:3726L21 siRNA (3708C) stab04	30700 HER2:3879U21 siRNA stab04		30702 HER2:2364L21 siRNA (2346C) stab05	30703 HER2:3708U21 siRNA stab05	HER2:3897L21 siRNA (3879C) stab05		HER2:3726L21 siRNA (3708C) stab08	30953 HER2:3708U21 siRNA stab04	HER2:3726L21 siRNA (3708C) stab05	31525 HRAS:77U21 siRNA	31526 HRAS:154U21 siRNA	31527 HRAS:459U21 siRNA	31528 HRAS:513U21 siRNA	31529 HRAS:95L21 siRNA (77C)
30448	30449	30645		30647	30648	30697	30698		30700	30701		30703	30704	30951	30952	30953	30954	31525	31526	31527	31528	31529
seuse	seuse	seuse	antisense	antisense	seuse	seuse	seuse	antisense	seuse	antisense	antisense	seuse	antisense	seuse	antisense	esues	antisense	sense	sense	seuse	sense	antisense
76	92	9/	75	92	75	78	76	75	79	78	92	75	79	75	75	22	75	80	81	82	83	80
GGUGCUUGGAU	GGUGCUUGGAU	4 GGUGCUUGGAUCUGGCGCU	3 UGGGGUCGUCAAAGACGUU															GAACCAUUUUGUGGACGAAUACG	GCCUGUUGGACAUCCUGGAUACC	GAGGAUGCCUUCUACACGUUGGU	CUGAACCCUCCUGAUGAGAGUGG	GAACCAUUUUGUGGACGAAUACG
		2344	3706	2344	3706			3706	3877	1882	2344	3706	3877	3706	3706	3706	e	_				95
Herz	Her2	Her2	Herz	Her2	Her2	Her2	Her2	Herz	Her2	Her2	Her2	Her2	Her2	Her2	Her2	Her2	Her2	HRAS	HRAS	HRAS	HKAS	HKAS

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290	591	292	593	594	595	296	297	298	299	009	601	602	603	604	909	909	209	809	609	610	611	612	613
UAUCCAGGAUGUCCAACAGTT	CAACGUGUAGAAGGCAUCCTT	ACUCUCAUCAGGAGGGUUCTT	AGCUUGGCCAAUCCGUGCGGU	UUGCGGAGGGUGGGCCUGGGA	CUGCCGCCUUCCACGCUCAU	ACCCACUGCCACCGCGAAGAG	GCGCCCGAUUCCCUGAGCUG	CGCACGGAUUGGCCAAGCUGA	CCAGGCCCACCCUCCGCAACC	GAACGGUGGAAGGCGGCAGGC	cunceceeueeckeueeeuec	GCUCAGGGAAUCGCGCGCGC	B ucccuuuAuAAGccGAcucTT B	B uuccAccGuucAuucuAGATT B	B ccAccGuucAuucuAGAGcTT B	B GAAGAGuuGGCcucuGucATT B	GAGucGGcuuAuAAAGGGATsT	ucuAGAAuGAAcGGuGGAATsT	GcucuAGAAuGAAcGGuGGTsT	uGAcAGAGcccAAcucuucTsT	B GAAGAGccAAcuGuGuGAGTT B	B AGGGAGGAGAGGAGuuccTT B	B GGAGUAcAGcAAAcuGAAGTT B
31530 HRAS:172L21 siRNA (154C)	31531 HRAS:477L21 siRNA (459C)	31532 HRAS:531L21 siRNA (513C)	29950 hTR:33U21 siRNA	29951 hTR:101U21 siRNA	29952 hTR:235U21 siRNA	29953 hTR:382U21 siRNA	29954 hTR:494U21 siRNA	29955 hTR:53L21 siRNA (33C)	29956 hTR:121L21 siRNA (101C)	29957 hTR:255L21 siRNA (235C)	29958 hTR:402L21 siRNA (382C)	29959 hTR:514L21 siRNA (494C)	30913 hTR:64U21 siRNA stab04	30914 hTR:243U21 siRNA stab04	30915 hTR:245U21 siRNA stab04	30916 hTR:397U21 siRNA stab04	30917 hTR:82L21 siRNA (64C) stab05	30918 hTR:261L21 siRNA (243C) stab05	30919 hTR:263L21 siRNA (245C) stab05		30801 IKKg:166U21 siRNA stab04	30802 IKKg:407U21 sIRNA stab04	30803 IKKg:1162U21 siRNA stab04
3153			2995	2995	2995	2995	2995	-		2995		2995	3091	3091	3091	3091		3091		30920	3080	3080	3080
antisense	antisense	antisense	sense	seuse	seuse	seuse	sense	antisense	antisense	antisense	antisense	antisense	seuse	seuse	esues	seuse	antisense	antisense	antisense	antisense	esues	seuse	sense
8	82	83	84	82	98	87	88	84	98	98	87	88	83	06	91	95	68	06	91	92	93	98	92
GCCUGUUGGACAUC	GAGGAUGCCUUCUACACGUUGGU	CUGAACCCUCCUGAUGAGAGUGG	UCAGCUUGGCCAAUCCGUGCGGU	GGUUGCGGAGGGU	_	GCACCCACUGCCAC	GCGCGCGCGAUUC	UCAGCUUGGCCAAUCCGUGCGGU	GGUUGCGGAGGGUGGGCCUGGGA	GCCUGCCGCCUCCACCGUUCAU	GCACCCACUGCCACCGCGAAGAG		GCUCCCUUNANAAGCCGACUCGC	ccuuccacceuucauucuagagc	UUCCACCGUUCAUUCUAGAGCAA	GCGAAGAGUUGGGCUCUGUCAGC	GCUCCCUUUANAAGCCGACUCGC	ccuuccacceuuca	UUCCACCGUUCAUUCUAGAGCAA	GCGAAGAGUUGGGCUCUGUCAGC	UGGAAGAGCCAACUGUGUGAGAU	AGAGGGAGGAGGAGUUCCUC	AGGGAGUACAGCAAACUGAAGGC
172		531	31	66	233	380	492	31	66	233	380	492	62	241	243	395	62	241	243	395	166	407	1162
HRAS	HRAS	HRAS	hTR	h R	hTR	hTR	hTR	hTR	hTR	hTR	hTR	hTR	hTR	hTR	hTR	hTR	hTR	hTR	hTR	hTR	IKKg	IKKg	IKKg

ST 615	sT 616		ST 617																						
B cAuGGAGuGGAuuGAGuAGTT	GGAAcuccuucuccuccuTsT		cuucAGuuuGcuGuAcuccTsT	cuucAGuuuGeuGuAcuccTsT	cuucAGuuuGcuGuAcuccTsT cuAcucAAuGcAcuccAuGTsT B AccucAAcuccuGccAcAATT B	cuucAGuuuGcuGuAcuccTsT cuAcucAAuGcAcuccAuGTsT B AccucAAcuccuGccAcAATT B B cuccuGucuGcAuuGcAcTT B	cuucAGuuuGeuGuAcuccTsT cuAcucAAuGcAcuccAuGTsT B AccucAAcuccuGccAcAATT B B cuccuGucuuGcAuuGcAcTT B B uuGcAcuuGucAAAAcAGTT B	cuucAGuuuGcuGuAcuccTsT cuAcucAAuGcAcuccAuGTsT B AccucAAcuccuGccAcAATT B B cuccuGucuuGcAuuGcAcTT B B uuGcAcuuGucAAAcAACTT B B cAcAGcuACAACAACTT B	cuucAGuuuGeuGuAcuccTsT cuAcucAAuGcAcuccAuGTsT B AccucAAcuccuGccAcAATT B B cuccuGucuuGcAuuGcAcTT B B uuGcAcuuGucAcAAAcAGTT B B cAcAGcuAcAAcaGTT B B cAcAGcuAcAAcaGATT B	cuucAGuuuGcuGuAcuccTsT cuAcucAAuGcAcuccAuGTsT B AccucAcuccuGccAcaCATT B B cuccuGucuGcAcuGcAcTT B B uuGcAcuuGucAcaAAcAGTT B B cAcAGcuAcAAcaAGTT B B cAcAGCuAcAAcaGGTT B GuGcAAuGcAAGAAGGTST	cuucAGuuuGcuGuAcuccTsT cuAcucAAuGcAcuccAuGTsT B AccucAAcuccuGccAcAATT E B cuccuGucuuGcAuuGcACTT B cuCaGcuAcAAcuGGAGTT B cAcAGcuAcAAcuGGAGATT CuGuGGCAGGAGuuGAGGUTs cuGuuuGuGAAAGAGGTs	cuucAGuuuGcuGuAcuccTsT cuAcucAAuGcAcuccAuGTsT B AccucAAcuccuGccAcAATT E B uuGcAcuuGucuuGcAuuGcAcTT E B uAcAGcuACACACGAGCATT I B uAcAGcuACAACGAGAGCATT I UuGuGGCAGGAGUUGAGGATS cuGuuuGuGACAAGGAGTS uGcuccAGuuGAAGGAGTS	cuucAGuuuGcuGuAcuccTsT cuAcucAAcuCcAcuCGCACAATT B B AccuCACUCGUCCACAATT B B cuccuGucuuGcAuuGcAcTT B B uuGcAcuUGuCACAACAGTT B B uuGcAcuUGCACAACAGTT B CuGGCAGGGAGUGAGGATT B cuGGCAGGGAGUGAGGCATT cuGuuuGuGACAAGAGGGTST aGCUCCAACUCCCCCACAATT ACCUCAACUCCCCCACAATT	cuucAGuuuGeuGuAcuccTsT cuAcucAAcuccauGcAAATT B B AccucAAcuccauGcACAATT B B cuccuGucuuGcAuuGcACTT B B uuGcAcuuGucAACaCAATT B B uuGcAcuuGucAACaCAATT B B cAcAGcuAcAACaCAACTT B CuGuuuGuCAACACACACTST cuGuuuGuCAACACACAATT CCUCAACUCCACCAATT CCUCAACUCCACCAATT CCUCAACUCCACCAATT CCUCCACCUCCACATT	cuucAGuuuGcuGuAcuccTsT cuAcucAAcuCcAcuCGCACAATT B B AccuCAACUCCUGCACAATT B B CUCCUGCACAACACTT B B UUGCACUCUUGCACUGCACATT B B UUGCAAUGCACAACACTT B CUCCUCAACUCCUGCCACAATT CUCCUCAACUCCACAATT CUCCACUCCAACUCCACAATT CUCCUCAACUCCACAATT CUCCUCAACUCCACAATT CUCCUCAACUCCACAATT CUCCUCAACUCCACAATT CUCCUCAACUCCACAATT CUCCUCAACUCCACAATT CUCCUCAACUCCACAATT CUCCUCAACUCCACAATT	cuucAGuuuGcuGuAcuccTsT cuAcucAAuGcAcucGuGTsT B AccucAAcuccuGcAcATT B B CACGCUACAACTT B B UUGCACUUGCAACAGTT B B CACGCUACAACGGGGGTT T UUGCAAUGGACAGGGTST CUGUUUGUGACAAGGGTST CUGUUUGUGACUCCCCCCCAATT CUCCUCAACUCCUGCCACATT CUCCUGUCCUGCCAACTT CACAGCUACAACGGGGGTT CACAGCUACAACUGCAACGTT CACAGCUACAACUGCAACGTT CACAGCUACAACUGCAACATT CACAGCUACAACUGCAACGTT CACAGCUACAACUGCAACATT CACAGCUACAACUGCAACATT CACAGCUACAACUGCAACACT CACAGCUACAACUGGAACACT CACAGCUACAACUGCAACACT CACAGCUACAACUGGAACACT CACAGCUACAACUCGAACACT CACAGCUACAACUCGAACACT CACAGCUACAACUCGAACACT CACAGCUACAACUCGAACACT CACAGCUACAACUCGAACACT CACAGCUACAACUCGAACACT CACAGCUACAACUCGAACACT CACAGCUACAACACACT CACAGCUACAACACACT CACAGCUACAACACACT CACAGCUACAACACACACT CACAGCUACAACACACT CACAGCUACAACACACACT CACAGCAACACACACACACT CACAGCAACACACACACACACACACACACACACACACAC	cuucAGuuuGcuGuAcuccTsT cuAcucAAuGcAcuccAuGTsT B AccucAAcuccuGccAcAATT B B CACGCUACACACATT B B UUGCACUUGCCAAACAGTT B B CACGCUACAACAGAGTT B UUCUGCACUCGAGACAGTT TS CUGUUUGUGACAGAGATST CUCUCAACUCCACAATT CUCCUGUCAACUGCCACATT CACAGCUACAACUGCCACAATT CACAGCUACAACUGCCACAATT CACAGCUACAACUGCCACAATT CACAGCUACAACUGCCACAATT CACAGCUACAACUGCCACAATT CACAGCUACAACUGCCACATT CACAGCUACAACUGCAACTT CACAGCUACAACUGCACATT CACAGCUACAACUGCACATT CACAGCUACAACUGCACATT CACAGCUACAACUGCACATT CACAGCUACAACUGCACATT CACAGCUACAACUGCACATT CACAGCUACAACUGCACATT CACAGCUACAACUGCACATT CACAGCUACAACUGAGGCATT CACAGCCAGGAGUUGAGGUTT	cuucAGuuuGeuGuAcuccTsT cuAcucAAuGeAcucGuGTsT B AccucAAcuccuGcAAATT B B CACGCUACAACAAACATT B B UUGCACUUGCAAACAGTT B B CACGCUACAACAGATT B B CACGCUACAACAGATT B CUGUUUGUGAACAGAGTST CUGUUUGUGACAAGACATT CACCAACUCCAACTT CACAACUCCAACTT CACAACACTAACAACT CUCCUGUCUUGCAATT CACAACTAACAACT CUCCUGUCUUGCAATT CACAACTAACAACT CUCCUGUCUGCCAACTT CACAGCUACAACTAACAATT CACAGCUACAACTAACAATT 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KRAS2	938	UUUCCUCGAAGUGCCAGUAUUCC	109	antisense	31538	31538 KRAS2:938L21 siRNA (920C)	AAUACUGGCACUUCGAGGATT	662
KRAS2	938	UNUCCUCGAAGUGCCAGUAUUCC	109	antisense	31538	31538 KRAS2:938L21 siRNA (920C)	AAUACUGGCACUUCGAGGATT	662
KRAS2	1017	KRAS2 1017 AUUUCUGUCUUGGGGUUUUUGGU	110	antisense	31539	31539 KRAS2:1017L21 siRNA (999C)	CAAAAACCCCAAGACAGAATT	663
KRAS2	1017	KRAS2 1017 AUUUCUGUCUUGGGGUUUUUGGU	110	antisense	31539	31539 KRAS2:1017L21 siRNA (999C)	CAAAAACCCCAAGACAGAATT	663
KRAS2	1031	GUUUUUGGUGCAUGCAGUUGAUU	Ξ	antisense	31540	31540 KRAS2:1031L21 sIRNA (1013C)	UCAACUGCAUGCACCAAAATT	664
KRAS2	1031	GUUUUGGUGCAUGCAGUUGAUU	=	antisense	31540	31540 KRAS2:1031L21 siRNA (1013C)	UCAACUGCAUGCACCAAAATT	664
MAPK1			112	seuse	30817	30817 MAPK1:424U21 siRNA stab04	B cAGAccuAcuGccAGAGAATT B	665
MAPK1	778	AUCACACAGGGUUCCUGACAGAA	113	seuse	30818	30818 MAPK1:778U21 siRNA stab04	B cAcAcAGGGuuccuGAcAGTT B	999
MAPK1	1718	1718 UUGGCUCUAGUCACUGGCAUCUC	114	seuse	30819	30819 MAPK1:1718U21 sIRNA stab04	B GGcucuAGucAcuGGcAucTT B	299
MAPK1	2525	2525 ACUGUGGAGUUGACUCGGUGUUC	115	seuse	30820	30820 MAPK1:2525U21 siRNA stab04	B uGuGGAGuuGAcucGGuGuTT B	899
	442	ACCAGACCUACUGCCAGAGAACC	112	antisense	30821	30821 MAPK1:442L21 siRNA (424C) stab05	uucucuGGcAGuAGGucuGTsT	699
	962	AUCACACAGGGUUCCUGACAGAA	113	antisense	30822	30822 MAPK1:796L21 siRNA (778C) stab05	cuGucAGGAAcccuGuGuGTsT	670
MAPK1	1736	1736 UUGGCUCUAGUCACUGGCAUCUC	114	antisense	30823	MAPK1:1736L21 siRNA (1718C) stab05	GAuGccAGuGAcuAGAGccTsT	671
MAPK1	2543	MAPK1 2543 ACUGUGGAGUUGACUCGGUGUUC	115	antisense	30824	MAPK1:2543L21 siRNA (2525C) stab05	AcAccGAGucAAcuccAcATsT	672
MAPK1 4	1280	MAPK1 1280 GCCUACUUGCUCAGUACCACGA	116	seuse	31586	31586 MAPK14:1280U21 siRNA	CUACUUUGCUCAGUACCACTT	673
MAPK1	1611	MAPK1 1811 UGUCUGUCUUUGUGGGAGGGUAA	117	seuse	31587	31587 MAPK14:1611U21 siRNA	UCUGUCUUUGUGGGAGGGUTT	674
MAPK1 4	2884	MAPK1 2884 AAAAGGGUCUUCUUGGCAGCUUA	118	sense	31588	31588 MAPK14:2884U21 siRNA	AAGGGUCUUCUUGGCAGCUTT	675
MAPK1 4	3556	MAPK1 3556 GGACUCUAAGCUGGAGCUCUUGG	119	sense	31589	31589 MAPK14:3556U21 siRNA	ACUCUAAGCUGGAGCUCUUTT	929
MAPK1 4	1298	MAPK1 1298 GCCUACUUGCUCAGUACCACGA 4	116	antisense	31590	MAPK14:1298L21 siRNA (1280C)	GUGGUACUGAGCAAAGUAGTT	677
MAPK1 4	1629	MAPK1 1629 UGUCUGUCUUUGUGGGAGGGUAA	117	antisense	31591	MAPK14:1629L21 siRNA (1611C)	ACCCUCCCACAAAGACAGATT	678
MAPK1	2902	2902 AAAAGGGUCUUCUUGGCAGCUUA	118	antisense	31592	31592 MAPK14:2902L21 siRNA	AGCUGCCAAGAAGACCCUUTT	629

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725	726	727	728	729	730	731	732	733	734	735	736	737	738	739	740	741	742	743	744	745
B AGuucuGAGucGGuuccAATT B	uuGGAAccGAcucAGAAcuTsT	B GuucAGuGucucuccAAAATT B	B uuuGcAGAuAGccuuGAGcTT B	B uccuGcuGcuucAuuGAcTT B	B GAcuGccAuGuGuucAucATT B	uuuuGGAGAGAcAcuGAAcTsT	GcucAAGGcuAucuGcAAATsT	GucAAuGAAAGcAGcAGGATsT	uGAuGAAcAcAuGGcAGucTsT	CUGCAGUACCUCUACCUGCTT	CGUCUCCUACUGCACCAGATT	UGGAGCCUGGAAGACCAGCTT	GUGACUCAGAAGGCUCAGGTT	GCAGGUAGAGGUACUGCAGTT	UCUGGUGCAGUAGGAGACGTT	GCUGGUCUUCCAGGCUCCATT	ccugaeccuucugaeucactt	B uGcAcGuAuAuGccGAGAuTT B	B ccAuAuuGGAGAuGcuGuuTT B	B AuuGcGCAuAuGCGAcAcuTT B
31395 MYC:1971U21 siRNA inv stab07	MYC:1989L21 siRNA (1971C) inv stab11	30833 Nogo:1043U21 siRNA stab04	30834 Nogo:1407U21 siRNA stab04	30835 Nogo:3211U21 siRNA stab04	30836 Nogo:3883U21 siRNA stab04		30838 Nogo:1425L21 siRNA (1407C) stab05	30839 Nogo:3229L21 siRNA (3211C) stab05	30840 Nogo:3901L21 siRNA (3883C) stab05	NogoR:512U21 siRNA	31058 NogoR:662U21 siRNA	31059 NogoR:1086U21 siRNA	31060 NogoR:1371U21 siRNA	31133 NogoR:530L21 siRNA (512C)	antisense 31134 NogoR:680L21 siRNA (662C)	31135 NogoR:1104L21 siRNA (1086C)	31136 NogoR:1389L21 siRNA (1371C)	PCNA:550U21 siRNA stab04	30842 PCNA:574U21 sIRNA stab04	30844 PCNA:839U21 siRNA stab04
31395	31398	3083	3083	3083	30836	30837		30838	30840	31057	31058	3105	3106		31134			30841	30842	3084
seuse	antisense	seuse	seuse	seuse	seuse	antisense	antisense	antisense	antisense	seuse	seuse	seuse	seuse	antisense	antisense	antisense	antisense	seuse	seuse	sense
131	131	132	133	134	135	132	133	134	135	136	137	138	139	136	137	138	139	140	141	142
ACAACCUUGGCUGAGUCUUGAGA	ACAACCUUGGCUGAGUCUUGAGA	ucenucaeueucucuccaaaaec		AUUCCUGCUGCUUCAUUGACAG	UNGACUGCCAUGUGUUCAUCAUC	1061 UCGUUCAGUGUCUCUCCAAAAGC	GUUUUGCAGAUAGCCUUGAGCAA	AUUCCUGCUGCUUCAUUGACAG	UNGACUGCCAUGUGUUCAUCAUC	cconecaeuaccucuaccuecae		ACUGGAGCCUGGAAGACCAGCUU		CCCUGCAGUACCUCUACCUGCAG	ACCGUCUCCUACUGCACCAGAAC	ACUGGAGCCUGGAAGACCAGCUU	UGGUGACUCAGAGGCUCAGGUG	UUUGCACGUAUAUGCCGAGAUCU		AAAUUGCGGAUAUGGGACACUUA
1969	1969	1043	1407	3211	3883	1061	1425	3229	3901	510	099	1084	1369	510	099	1084	1369	548	225	837
MYC	MYC	Nogo	Nogo	Nogo	Nogo	Nogo	Nogo	Nogo	Nogo	NOGO R	NOGO R	NOGO R	NOGO R	NOG R	NOGO R	NOGO R	NOGO R	PCNA	PCNA	PCNA

746	747	748	749	750	751	752	753	754	755	756	757	758	759	760	761	761	762	762	762	762	763	763
AucucGGcAuAuAcGuGcATsT	AAcAGcAucuccAAuAuGGTsT	AGuGuccAuAuccGcAAuTsT	UGCACGUAUAUGCCGAGAUTT	CCAUAUUGGAGAUGCUGUUTT	AAAGCCACUCCACUCUUTT	AUUGCGGAUAUGGGACACUTT	AUCUCGGCAUAUACGUGCATT	AACAGCAUCUCCAAUAUGGTT	AAGAGAGUGGAGUGGCUUUTT	AGUGUCCCAUAUCCGCAAUTT	B AAAGccAcuccAcucucuuTT B	AAGAGAGAGAGAGCCCUUTST	B uucucAccucAccGAAATT B	uuucGGuGAGGuGAGAATsT	B cAGGAccuccAcAuGAuAGTT B	B cAGGAccuccAcAuGAuAGTT B	B AGAuuuGAccuuccuGAcATT B	B AGAuuuGAccuuccuGAcATT B	B AGAuuuGAccuuccuGAcATT B	B AGAuuuGAccuuccuGAcATT B	B uGAGuAGcuGGAuuAcAGGTT B	B uGAGuAGcuGGAuuAcAGGTT B
antisense 30845 PCNA:568L21 siRNA (550C) stab05	antisense 30846 PCNA:592L21 siRNA (574C) stab05	30848 PCNA:857L21 siRNA (839C) stab05	31033 PCNA:550U21 siRNA	31034 PCNA:574U21 siRNA	31035 PCNA:767U21 siRNA	31036 PCNA:839U21 siRNA	31109 PCNA:568L21 siRNA (550C)	31110 PCNA:592L21 siRNA (574C)	31111 PCNA:785L21 siRNA (767C)	31112 PCNA:857L21 siRNA (839C)	31310 PCNA:767U21 siRNA stab04	31311 PCNA:785L21 siRNA (767C) stab05	31322 PCNA:767U21 siRNA inv stab04	PCNA:785L21 siRNA (767C) inv stab05	30969 PKR:533U21 siRNA stab04	PKR:533U21 siRNA stab04	30970 PKR:1171U21 siRNA stab04	30970 PKR:1171U21 siRNA stab04	30970 PKR:1171U21 siRNA stab04	PKR:1171U21 siRNA stab04	30971 PKR:2430U21 siRNA stab04	PKR:2430U21 sIRNA stab04
30845	30846		31033	31034	31035	31036	31109				31310	31311	31322	31323	30969	30969	30970	30970	30970	30970	30971	30971
antisense	antisense	antisense	seuse	sense	seuse	sense	antisense	antisense	antisense	antisense	seuse	antisense	seuse	antisense	seuse	seuse	seuse	seuse	seuse	seuse	seuse	sense
140	141	142	140	141	143	142	140	141	143	142	143	143	143	143	144	144	25	22	22	22	22	22
UUUGCACGUAUAUGCCGAGAUCU	AGCCAUAUUGGAGAUGCUGUUGU	AAAUUGCGGAUAUGGGACACUUA	UNUGCACGUANAUGCCGAGAUCU	AGCCAUAUUGGAGAUGCUGUUGU	CAAAAGCCACUCCACUCUCUCA	AAAUUGCGGAUAUGGGACACUUA	UUUGCACGUAUAUGCCGAGAUCU	AGCCAUAUUGGAGAUGCUGUUGU	CAAAAGCCACUCCACUCUUCA	AAAUUGCGGAUAUGGGACACUUA	CAAAAGCCACUCCACUCUCA	CAAAAGCCACUCCACUCUCA	CAAAAGCCACUCCACUCUUCA	CAAAAGCCACUCCACUCUUCA	UUCAGGACCUCCACAUGAUAGGA	UUCAGGACCUCCACAUGAUAGGA	AACAACCACAAAAUACAACAAGA	AACAACCACAAAAUACAACAAGA	AACAACCACAAAAUACAACAAGA	AACAACCACAAAAUACAACAAGA	AACAACCACAAAAUACAACAAGA	AACAACCACAAAAUACAACAAGA
548	572	837	548	572	765	837	548	572	765	837	765	765	765	765	533	533	1171	1171	1171	1171	2430	2430
PCNA	PCNA	PCNA	PCNA	PCNA	PCNA	PCNA	PCNA	PCNA	PCNA	PCNA	PCNA	PCNA	PCNA	PCNA	PKR	PKR	PKR	PKR	PKR	PKR	PKR	PKR

	764	765	765	299	766	792	767	768	768	769	770	77.1	772	773	774	775	776	111	778	779	780	781
	B GGucucAAcuccuGAccuTT B	cuAucAuGuGGAGGuccuGTsT	cuAucAuGuGGAGGuccuGTsT	uGucAGGAAGGucAAAucuTsT	uGucAGGAAGGucAAAucuTsT	ccuGuAAuccAGcuAcucATsT	ccuGuAAuccAGcuAcucATsT	AGGucAGGAGuuuGAGAccTsT	AGGucAGGAGuuuGAGAccTsT	B AAAGGcuGAGGuuGcuGAuTT B	B AAAcAAccuuccAAcAAccTT B	B AAGGAcuGAuGAccAAAcATT B	AucAGcAAccucAGccuuuTsT	GGuuGuuGGAAGGuuGuuuTsT	uGunuGGucAucAGuccuuTsT	AAAGGCUGAGGUUGCUGAUTT	AAACAACCUUCCAACAACCTT	GGAUGUGGUGAUUCAGGAUTT	AAGGACUGAUGACCAAACATT	AUCAGCAACCUCAGCCUUUTT	GGÜÜGÜÜGGAAGGÜÜGÜÜÜTT	AUCCUGAAUCACCACAUCCTT
sus/2 ren.23/802/sirving	30972 PKR:2518U21 siRNA stab04	30973 PKR:551L21 siRNA (533C) stab05	30973 PKR:551L21 siRNA (533C) stab05	30974 PKR:1189L21 siRNA (1171C) stab05	30974 PKR:1189L21 siRNA (1171C) stab05	30975 PKR:2448L21 siRNA (2430C) stab05		PKR:2536L21 siRNA (2518C) stab05	30976 PKR:2536L21 siRNA (2518C) stab05	30713 PRKCA:519U21 siRNA stab04	30714 PRKCA:1000U21 sIRNA stab04	30716 PRKCA:1736U21 siRNA stab04	30717 PRKCA:537L21 siRNA (519C) stab05	PRKCA:1018L21 siRNA (1000C) stab05	PRKCA:1754L21 siRNA (1736C) stab05	30989 PRKCA:519U21 siRNA	30990 PRKCA:1000U21 siRNA	30991 PRKCA:1143U21 siRNA	30992 PRKCA:1736U21 siRNA	31065 PRKCA:537L21 sIRNA (519C)	31066 PRKCA:1018L21 siRNA (1000C)	
.,,,,	30972	30973	30973	30974	30974	30975	30975	30976	30976	30713	30714	30716	30717	30718	30720	30989	30990	30991	30992			31067
201120	seuse	antisense	antisense	antisense	antisense	antisense	antisense	antisense	antisense	seuse	seuse	sense	antisense	antisense	antisense	sense	seuse	seuse	seuse	antisense	antisense	antisense 31067
Š	25	22	22	22	22	22	22	22	22	145	146	147	145	146	147	145	146	148	147	145	146	148
AACAACCACAAAAUACAACAAGA	AACAACCACAAAAUACAACAAGA	AACAACCACAAAAUACAACAAGA	AACAACCACAAAAUACAACAAGA	AACAACCACAAAAUACAACAAGA	AACAACCACAAAAUACAACAAGA	AACAACCACAAAAUACAACAAGA	AACAACCACAAAAUACAACAAGA	AACAACCACAAAAUACAACAAGA	AACAACCACAAAAUACAACAAGA	CUAAAGGCUGAGGUUGCUGAUGA	GGAAACAACCUUCCAACAACCUU	CAAAGGACUGAUGACCAAACACC	CUAAAGGCUGAGGUUGCUGAUGA	GGAAACAACCUUCCAACAACCUU	CAAAGGACUGAUGACCAAACACC	CUAAAGGCUGAGGUUGCUGAUGA	GGAAACAACCUUCCAACAACCUU	AAGGAUGUGGUGAUCAGGAUGA	CAAAGGACUGAUGACCAAACACC	CUAAAGGCUGAGGUUGCUGAUGA	GGAAACAACCUUCCAACAACCUU	AAGGAUGUGGUGAUCAGGAUGA
8167	2518	551	551	1189	1189	2448	2448	2536	2536	517	866	1734	517	866	1734	1		1141	1734	517	866	1141
Į X	PKR	PKR	PKR	PKR	PKR	PKR	PKR	PKR	PKR	PRKCA	PRKCA	PRKCA	PRKCA	PRKCA	PRKCA	PRKCA	PRKCA	PRKCA	PRKCA	PRKCA	PRKCA	PRKCA

782	783	784	785	786	787	788	789	790	791	792	793	794	795	796	797	798	799	800	801	802
UGUUUGGUCAÜCAGUCCUUTT	B GGAuGuGGuGAuucAGGAuTT B	AuccuGAAucAccAcAuccTsT	B GGAuGuGGuGAucAGGAuTT B	AuccuGAAucAccAcAuccTsT	B uAGGAcuuAGuGGuGuAGGTT B	ccuAcAccAcuAAGuccuATsT	B uAGGAcuuAGuGGuGuAGGTT B	ccuAcAccAcuAAGuccuATsT	CUCGUUCCUCUUGGACAAGTT	GGUGAGCUACAAACACAUGTT	AGUGGAAGACUGGCUGAGCTT	CCUCUAGCCUGUUGUUGUTT	CUUGUCCAAGAGAACGAGTT	CAUGUGUUGUAGCUCACCTT	GCUCAGCCAGUCUUCCACUTT	ACAACAACAGGCUAGAGGTT	B uccGAcAuGAAGccAGuGATT B	B cuGAuGGAcAAGAGGAAAGTT B	B GuGuGGAuAAGGcuuAGGuTT B	ucAcuGGcuucAuGucGGATsT
antisense 31068 PRKCA:1754L21 siRNA (1736C)	31376 PRKCA:1143U21 siRNA stab04	79 PRKCA:1161L21 siRNA (1143C) stab05	31382 PRKCA:1143U21 siRNA stab07	35 PRKCA:1161L21 siRNA (1143C) stab11	31388 PRKCA:1143U21 siRNA inv stab04		31394 PRKCA:1143U21 siRNA inv stab07	77 PRKCA:1161L21 siRNA (1143C) inv stab11	31557 PTP4A3:205U21 siRNA	31558 PTP4A3:367U21 siRNA	31559 PTP4A3:574U21 siRNA	31560 PTP4A3:1168U21 siRNA	31561 PTP4A3:223L21 siRNA (205C)	32 PTP4A3:385L21 siRNA (367C)	31563 PTP4A3:592L21 siRNA (574C)	4 PTP4A3:1186L21 siRNA (1168C)	30865 PTPN1:242U21 siRNA stab04	77 PTPN1:874U21 siRNA stab04	8 PTPN1:3037U21 siRNA stab04	i9 PTPN1:260L21 siRNA (242C) stab05
3106	3137	31379	3138	31385	3138	31391	3139	31397	3155	3155	3155	3156	-	31562	_	31564	3086	30867	30868	3086
antisense	sense	antisense	seuse	antisense	sense	antisense	seuse	antisense	seuse	seuse	seuse	seuse	antisense	antisense	antisense	antisense	seuse	seuse	esues	antisense 30869
147	148	148	148	148	148	148	148	148	149	150	151	152	149	150	151	152	153	154	155	153
PRKCA 1734 CAAAGGACUGAUGACCAAACACC	AAGGAUGUGGUGAUUCAGGAUGA	AAGGAUGUGGUGAUUCAGGAUGA	PRKCA 1141 AAGGAUGUGGUGAUUCAGGAUGA	PRKCA 1141 AAGGAUGUGGUGAUCAGGAUGA	1141 AAGGAUGUGGUGAUUCAGGAUGA	AAGGAUGUGGUGAUUCAGGAUGA	PRKCA 1141 AAGGAUGUGGUGAUUCAGGAUGA			GAGGUGAGCUACA		cuccucuAeccuenuuenuenee		GAGGUGAGCUACAACACAUGCG	GUAGUGGAAGACUGGCUGAGCCU	1186 CUCCUCUAGCCUGUUGUUGUGG	NAUCCGACAUGAAGCCAGUGACU	UGCUGAUGGACAAGAGGAAAGAC	AGGUGUGGAUAAGGCUUAGGUGC	UAUCCGACAUGAAGCCAGUGACU
1734	1141	1141	1141	1141	1141	1141	1141	1141	205	367	574	1168		_	592	1186		872		240
PRKCA	PRKCA 1141	PRKCA	PRKCA	PRKCA	PRKCA	PRKCA	PRKCA	PRKCA	PTP4A 3	PTP4A 3	PTP4A 3	PTP4A 3	PTP4A 3	PTP4A 3	PTP4A 3		PTPN1	PTPN1	PTPN1	PTPN1

	830	831	832	833	834	835	836	837	838	839	840	841	842	843	844	845	846	847	848	849	850	851
	CUCAGCCUCAUAGAAGCCATT	CUUUCUGCACCUUGUCACATT	AGAGUACCAGAAGCUGGAGTT	B cuccAGcuncuGGuAcucuTT B	AGAGUACCAGAAGCUGGAGTST	B ucucAuGGucuucGAccucTT B	GAGGucGAAGAccAuGAGATsT {	B uAuGcuGuGGuGcuuAAuGTT B	B uGcuGGAcAuGAGAuGGAGTT B	B GAGGcuAcAGGGGuuAGccTT B	B GAccuAccucAAAGGGcAGTT B	CAUUAAGCACACACACAUATST	cuccAucucAuGuccAGcATsT	GGcuAAcccouGuAGccucTsT	cuGcccuuuGAGGuAGGucTsT	UAUGCUGUGGUGCUUAAUGTT			GACCUACCUCAAAGGGCAGTT 8	CAUUAAGCACCACAGCAUATT	CUCCAUCUCAUGUCCAGCATT 8	GGCUAACCCCUGUAGCCUCTT 8
(146C)	31106 RelA:308L21 siRNA (290C)	31107 ReIA:663L21 siRNA (645C)	31108 ReIA:1975L21 siRNA (1957C)	31308 ReIA:1957U21 siRNA stab04	31309 RelA:1975L21 siRNA (1957C) stab05	31320 RELA:1957U21 siRNA inv stab04	31321 RELA:1975L21 sIRNA (1957C) inv stab05	30873 SCD:995U21 siRNA stab04	30874 SCD:2520U21 siRNA stab04	30875 SCD:3785U21 siRNA stab04	30876 SCD:4774U21 siRNA stab04	30877 SCD:1013L21 siRNA (995C) stab05		30879 SCD:3803L21 siRNA (3785C) stab05	30880 SCD:4792L21 siRNA (4774C) stab05	31021 SCD:995U21 siRNA	31022 SCD:2520U21 siRNA	31023 SCD:3785U21 sIRNA	31024 SCD:4774U21 siRNA	31097 SCD:1013L21 siRNA (995C)	31098 SCD:2538L21 siRNA (2520C)	antisense 31099 SCD:3803L21 siRNA
	antisense 31	antisense 31	antisense 31	sense 31	antisense 31	sense 31	antisense 31	sense 30	sense 30	sense 30	sense 30	antisense 30	antisense 30878	antisense 308	antisense 308	sense 310	sense 31(sense 31(sense 310	antisense 310	antisense 310	ntisense 310
	162	163	49	164	164	164	164	165	166	167	168	165 a	166	167 a	168	165	166	167	168	165 a	166 a	167 a
	GAUGGCUUCUAUGAGGCUGAGCU	UGUGUGACAAGGUGCAGAAGAG	uccuccaecuucueeuacucucc		uccuccaecuucueeuacucucc		nccuccaecuncueeuacuccc	GAUAUGCUGUGGUGCUUAAUGCC					ACUGCUGGACAUGAGAUGGAGAG	UAGAGGCUACAGGGGUUAGCCUG	CUGACCUACCUCAAAGGGCAGUU	GAUAUGCUGUGGUGCUUAAUGCC	ACUGCUGGACAUGAGAUGGAGAG	3783 UAGAGGCUACAGGGGUUAGCCUG		GAUAUGCUGUGGUGCUUAAUGCC	ACUGCUGGACAUGAGAUGGAGAG	UAGAGGCUACAGGGGUUAGCCUG
-		643	1955	-		1955	1955	663	2518	3783	4772	993	2518	3783	4772	993	2518	3783	_	993	2518	3783
i	KELA	RELA	RELA	RELA	RELA	RELA	RELA	SCD	SCD	SCD	SCD	SCD	SCD	SCD	SCD	SCD	SCD	SCD	SCD	SCD	SCD	SCD

SCD 47	4772	CUGACCUACCUCAAA	168	antisense	31100	antisense 31100 SCD:4792L21 siRNA (4774C)	CUGCCCUUUGAGGUAGGUCTT	852
	_	CUGCGCACGUGGGAAGCCCUGGC	169	seuse	29960		GCGCACGUGGGAAGCCCUGGC	853
2	-	UGCAGAGGCUGUGCGAGCGCGGC		sense	29961		CAGAGGCUGUGCGAGCGCGGC	854
إم	641		_	sense	29962		UCUGGGAUGCGAACGGGCCUG	855
	1244	CUUGGGAACCACGCGCAGUGCCC	172	sense	29963	TERT:1246U21 siRNA	UGGGAACCACGCGCAGUGCCC	856
5	495	2495 UGCCACCACGCGUGCGCAUCAG	173	sense	29964	TERT:2497U21 siRNA	CCACCACGCGUGCGCAUCAG	857
	-	CUGCGCACGUGGGAAGCCCUGGC		antisense	29965		CAGGGCUUCCCACGUGCGCAG	858
3		UGCAGAGGCUGUGCGAGCGCGGC	170	antisense	29966	TERT:331L21 sIRNA (311C)	CGCGCUCGCACAGCCUCUGCA	859
Ö		CGUCUGGGAUGCGAACGGGCCUG	171	antisense	29967	TERT:663L21 siRNA (643C)	GCCCGUUCGCAUCCCAGACG	860
7		CUUGGGAACCACGCGCAGUGCCC	172	antisense	29968	TERT:1266L21 siRNA (1246C)	GCACUGCGCGUGGUUCCCAAG	861
24		UGCCACCACGCCGUGCGCAUCAG	173	antisense	29969	TERT:2517L21 sIRNA (2497C)	GAUGCGCACGGCGUGGUGGCA	862
11	1136	GUGGAGACCAUCUUCUGGGUUC	174	sense	30905	TERT:1138U21 siRNA stab04	B GGAGAccAucuucuGGGuTT B	863
TERT 17		AGUGUCUGGAGCAAGUUGCAAAG	175	sense	30906	TERT:1792U21 siRNA stab04	B uGucuGGAGcAAGuuGcAATT B	864
8		AUCAGAGCCAGUCUCACCUUCAA	176	sense	30907	TERT:2917U21 siRNA stab04	B cAGAGccAGucucAccuucTT B	865
82	2994	UGAAGUGUCACAGCCUGUUUCUG	177	seuse	30908	TERT:2996U21 siRNA	B AAGuGucAcAGccuGuuucTT B	866
1	_	GUGGAGACCAUCUUCUGGGUUC	174	antisense	60608	TERT:1156L21 siRNA (1138C) stab05	AcccAGAAAGAuGGucuccTsT	867
17			175	antisense	30910	TERT:1810L21 siRNA (1792C) stab05	uuGcAAcuuGcuccAGAcATsT	868
82		AUCAGAGCCAGUCUCACCUUCAA	176	antisense	30911	TERT:2935L21 siRNA (2917C) stab05	GAAGGuGAGAcuGGcucuGTsT	869
		UGAAGUGUCACAGCCUGUUUCUG	177	antisense	30912	TERT:3014L21 siRNA (2996C) stab05	GAAAcAGGcuGuGAcAcuuTsT	870
		AGGGAUAACACACUGCAAGUGGA	178	seuse	30881	TGFb:1528U21 siRNA stab04	B GGAuAAcAcAcuGcAAGuGTT B	871
TGFB1 23(_	CCAUAGCAACACUCUGAGAUGGC	179	seuse	30882	30882 TGFb:2385U21 siRNA stab04	B AuAGcAAcacucuGAGAuGTT B	872
24		GAACCUGCUUNAGUGGGGGAUAG	180	seuse	30883	TGFb:2486U21 siRNA stab04	B AccuGcuuuAGuGGGGGAuTT B	873
25(UAGCACUUUUGGGAGGCAGAGAU	181	seuse	30884	TGFb:2568U21 siRNA stab04	B GcAcuuuuGGGAGGcAGAGTT B	874
5	1526	AGGGAUAACACACUGCAAGUGGA	178	antisense	30885	30885 TGFb:1546L21 siRNA	cAcuuGcAGuGuGuuAuccTsT	875

			L			(4538C) = 1=1-05		
TGFB1	2383	CCAUAGCAACACUCUGAGAUGGC	179	antisense	30886		cAucucAGAGuGuuGcuAuTsT	876
TGFR1	2484		-			(2385C) stab05) }
5			180	antisense	30887	TGFb:2504L21 siRNA (2486C) stab05	AuccecAcuAAAGcAGGuTsT	877
IGFB1	-	UAGCACUUUUGGGA	181	antisense	30888	TGFb:2586L21 siRNA (2568C) stab05	cucuGccucccAAAAGuGcTsT	878
TGFB1	_		Н	sense	31053	31053 TGFb:1528U21 siRNA	GGAUAACACACUGCAAGUGTT	879
GFB1	$\overline{}$		179	seuse	31054	31054 TGFb:2385U21 siRNA	AUAGCAACACUCUGAGAUGTT	880
GFB1			180	seuse	31055	31055 TGFb:2486U21 siRNA	ACCUGCUUNAGUGGGGGAUTT	881
TGFB1	_		181	sense	31056	31056 TGFb:2568U21 siRNA	GCACUUUUGGGAGGCAGAGTT	882
IGFB1		1	178	antisense	31129	TGFb:1546L21 siRNA (1528C)	CACUUGCAGUGUGUUAUCCTT	883
TGFB1			179	antisense		31130 TGFb:2403L21 siRNA (2385C)	CAUCUCAGAGUGUUGCUAUTT	884
TGFB1	2484		180	antisense 31131		TGFb:2504L21 siRNA (2486C)	AUCCCCACUAAAGCAGGUTT	885
TGFB1	2566	1	181	antisense	31132		CUCUGCCUCCCAAAAGUGCTT	988
TNF F		AAGGACACCAUGAGGCACUGAAAG	182	seuse	30889	TNFa:79U21 siRNA	B GGAcAccAuGAGcAcuGAATT B	887
TN TN		UNGUNCCUCAGCCUCUNCUCCUN	183	sense	30890	30890 TNFa:178U21 sIRNA stab04	B GuuccucAGccucuucuccTT B	888
	568	CUCCUACCAGACCAAGGUCAACC	184	seuse	30891	TNFa:570U21 siRNA stab04	B ccuAccAGAccAAGGucAATT B	889
	1150	UNAGGCCUUCCUCUCCAGAUG	185	seuse	30892	TNFa:1152U21 siRNA stab04	B AGGccuuccucuccAGATT B	890
HN.		AAGGACACCAUGAGGCACUGAAAG	182	antisense	30893	TNFa:97L21 siRNA (79C) stab05	uucAGuGcucAuGGuGuccTsT	891
L L	\neg	UUGUUCCUCAGCCUCUUCUCCUU	183	antisense 30894		TNFa:196L21 siRNA (178C) stab05	GGAGAAGAGGCuGAGGAAcTsT	892
	568	CUCCUACCAGACCAAGGUCAACC	184	antisense 30895	30895	TNFa:588L21 siRNA (570C) stab05	uuGAccuuGGucuGGuAGGTsT	893
	1150	UNAGGCCUUCCUCUCCAGAUG	185	antisense	30896	TNFa:1170L21 siRNA	ucuGGAGAGGAAGGccuTsT	894
H H	-	AAGGACACCAUGAGCACUGAAAG	182	sense	31408	31408 TNFa:79U21 siRNA	GGACACCAUGAGCACUGAATT	895
	0/1	Significant Second Seco	183	sense	31409 1	31409 TNFa:178U21 siRNA	GUUCCUCAGCCUCUUCUCCTT	968
+	+	CUCCUACCAGACCAAGGUCAACC	184	1	31410	31410 TNFa:570U21 siRNA	CCUACCAGACCAAGGUCAATT	897
+	-	AACCACACACACACACACACACACACACACACACACAC	\neg	seuse	31411	31411 TNFa:1152U21 siRNA	AGGCCUUCCUCUCCAGATT	898
	1	ANGONCHOCHOCHCOCHCOCHAGE	182	antisense	31412	antisense 31412 TNFa:97L21 siRNA (79C)	UUCAGUGCUCAUGGUGUCCTT	668

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TNF	176	TNF 176 UUGUUCCUCAGCCUCUUCUCCUU	183	antisense 31	UCUUCUCCUU 183 antisense 31413 TNFa:196L21 siRNA (1780)	GGAGAAGAGGCUGAGGAACTT	006
TN.	268	TNF 568 CUCCUACCAGACCAAGGUCAACC	184	antisense 31	AAGGUCAACC 184 antisense 31414 TNFa:588L21 siRNA	UUGACCUUGGUCUGGUAGGTT	901
TNF	1150	TNF 1150 UVAGGCCUUCCUCUCCCAGAUG	185	antisense 31	UCUCCAGAUG 185 antisense 31415 TNFa:1170L21 siRNA	UCUGGAGAGGAAGGCCUTT	902
					(1152C)		_

Uppercase = ribonucleotide u,c = 2'-deoxy-2'-fluoro U,C T = thymidine B = inverted deoxy abasic s = phosphorothioate linkage A = deoxy Adenosine G = deoxy Guanosine

Table II

A. 2.5 μmol Synthesis Cycle ABI 394 Instrument

Reagent	Equivalents	Amount	Wait Time* DNA	Wait Time* 2'-O-methyl	Wait Time*RNA
Phosphoramidites	6.5	163 µL	45 sec	2.5 min	7.5 min
S-Ethyl Tetrazole	23.8	238 µL	45 sec	2.5 min	7.5 min
Acetic Anhydride	100	233 µL	5 sec	5 sec	5 sec
N-Methyl Imidazole	186	233 µL	5 sec	5 sec	5 sec
TCA	176	2.3 mL	21 sec	21 sec	21 sec
lodine	11.2	1.7 mL	.45 sec	45 sec	45 sec
Beaucage	12.9	645 µL	100 sec	300 sec	300 sec
Acetonitrile	NA	6.67 mL	NA	NA	NA NA

B. 0.2 µmol Synthesis Cycle ABI 394 Instrument

Reagent	Equivalents	Amount	Wait Time* DNA	Wait Time* 2'-O-methyl	Wait Time*RNA
Phosphoramidites	15	31 µL	45 sec	233 sec	465 sec
S-Ethyl Tetrazole	38.7	31 µL	45 sec	233 min	465 sec
Acetic Anhydride	655	124 µL	5 sec	5 sec	5 sec
N-Methyl Imidazole	1245	124 µL	5 sec	5 sec	5 sec
TCA	700	732 µL	10 sec	10 sec	10 sec
lodine	20.6	244 µL	15 sec	15 sec	15 sec
Beaucage	7.7	232 µL	100 sec	300 sec	300 sec
Acetonitrile	NA	2.64 mL	NA	NA	NA NA

C. 0.2 µmol Synthesis Cycle 96 well Instrument

Reagent	Equivalents:DNA/ 2'-O-methyl/Ribo	Amount: DNA/2'-O- methyl/Ribo	Wait Time* DNA	Wait Time* 2'-O- methyl	Wait Time* Ribo
Phosphoramidites	22/33/66	40/60/120 μL	60 sec	180 sec	360sec
S-Ethyl Tetrazole	70/105/210	40/60/120 μL	60 sec	180 min	360 sec
Acetic Anhydride	265/265/265	50/50/50 μL	10 sec	10 sec	10 sec
N-Methyl Imidazole	502/502/502	50/50/50 μL ·	10 sec	10 sec	10 sec
TCA	238/475/475	250/500/500 µL	15 sec	15 sec	15 sec
lodine	6.8/6.8/6.8	80/80/80 µL	30 sec	30 sec	30 sec
Beaucage	34/51/51	80/120/120	100 sec	200 sec	200 sec
Acetonitrile	NA	1150/1150/1150 µL	NA	NA	NA NA

- Wait time does not include contact time during delivery.
- Tandem synthesis utilizes double coupling of linker molecule

Table III

Group	Solution on	Stock VEGF	Number	Injectate	Dose	Conc.
	Filter (1.0	concentration	of	. (6.0 μL)		injectate
	μL)		Animals			
1 WEST	n 14/4 midding	NTA		Carrier Control	1/20 N (3/20)	
1	Tris-Cl pH	NA	5	water	NA	NA
	6.9	0.50 / 7				
2	R&D Systems	3.53 μg/μL	5	water	NA	NA
	VEGF-carrier]
	free					
	75 μM	0.50 / 7				
3	R&D Systems	3.53 μg/μL	5	Site 2340	10	1.67
	VEGF-carrier			Stab1	μg/eye	μg/μL
	free			siRNA		
	75 μM	0.50 / 7		22.12		
4	R&D Systems	3.53 μg/μL	5	Site 2340	3	0.5
	VEGF-carrier			Stab1	μg/eye	μg/μĽ
	free			siRNA		
5	75 μM	0 F0 / . T	r-	6: 0040		0.167
	R&D Systems VEGF-carrier	3.53 μg/μL	5	Site 2340	1	0.167
	free			Stab1 siRNA	μg/eye	, µg/µĽ
1	75 μM			SIKIVA		
6	R&D Systems	3.53 µg/µL	5	Inactive	10	1.67
	VEGF-carrier	3.33 μg/ μL		Site 2340		i
	free			Stab1	μg/eye	μg/μL
	75 μM			siRNA		
7	R&D Systems	3.53 µg/µL	. 5	Inactive	3	0.5
'	VEGF-carrier	0.00 μg/ μL	. 3	Site 2340	_	
	free			Stab1	μg/eye	μg/μL
	75 μM			siRNA		
8	R&D Systems	3.53 µg/µL	5	Inactive	1	0.167
	VEGF-carrier	0.00 µ6/ µL	J	Site 2340	μg/eye	0.107 μg/μL
	free			Stab1	- 146/ C) C	μ ₆ / μυ
	75 μM			siRNA		

Table IV

Non-limiting examples of Stabilization Chemistries for chemically modified siNA constructs

Chemistry	pyrimidine	Purine	cap	p=S	Strand
"Stab 1"	Ribo	Ribo	-	5 at 5'-end 1 at 3'-end	S/AS
"Stab 2"	Ribo	Ribo	-	All linkages	Usually AS
"Stab 3"	2'-fluoro	Ribo	-	4 at 5'-end 4 at 3'-end	Usually S
"Stab 4"	2'-fluoro	Ribo	5' and 3'- ends	-	Usually S
"Stab 5"	2'-fluoro	Ribo	-	1 at 3'-end	Usually AS
"Stab 6"	2'-O-Methyl	Ribo	5' and 3'- ends	-	Usually S
"Stab 7"	2'-fluoro	2'-deoxy	5' and 3'- ends	-	Usually S
"Stab 8"	2'-fluoro	2'-O- Methyl	-	1 at 3'-end	Usually AS
"Stab 9"	Ribo	Ribo	5' and 3'- ends	-	Usually S
"Stab 10"	Ribo	Ribo	_	1 at 3'-end	Usually AS
"Stab 11"	2'-fluoro	2'-deoxy	-	1 at 3'-end	Usually AS

⁵ CAP = any terminal cap, see for example Figure 10.

All Stab 1-11 chemistries can comprise 3'-terminal thymidine (TT) residues

All Stab 1-11 chemistries typically comprise 21 nucleotides, but can vary as described herein.

S = sense strand

10 AS = antisense strand

Table V

	Table V
Acc#	Description
NM_002825	Homo sapiens pleiotrophin (heparin binding growth factor 8, neurite growth-
	promoting factor 1) (PTN), mRNA
NM_033418	Homo sapiens hypothetical protein MGC9084 (MGC9084), mRNA
NM_033111	Homo sapiens LOC88523 (LOC88523), mRNA
NM_032564	Homo sapiens diacylglycerol O-acyltransferase homolog 2 (mouse) (DGAT2),
	mRNA
NM_032311	Homo sapiens KIAA1649 protein (KIAA1649), mRNA
NM_022130	Homo sapiens golgi phosphoprotein 3 (coat-protein) (GOLPH3), mRNA
NM_021980	Homo sapiens optineurin (OPTN), mRNA
NM_000660	Homo sapiens transforming growth factor, beta 1 (Camurati-Engelmann disease) (TGFB1), mRNA
NM_020423	Homo sapiens hypothetical protein LOC57147 (LOC57147), mRNA
NM_020351	Homo sapiens smooth muscle cell-expressed and macrophage conditioned medium-induced protein smag-64 (LOC57086), mRNA
NM 019556	Homo sapiens hypothetical protein dJ473B4 (DJ473B4), mRNA
NM_018676	Homo sapiens TMTSP for transmembrane molecule with thrombospondin
ن –	module (LOC55901), mRNA
NM_016265	Homo sapiens GIOT-3 for gonadotropin inducible transcription repressor-3
	(GIOT-3), mRNA
NM_016531	Homo sapiens Kruppel-like factor 3 (basic) (KLF3), mRNA
NM_016372	Homo sapiens seven transmembrane domain orphan receptor (TPRA40), mRNA
NM_016211	Homo sapiens yeast Sec31p homolog (KIAA0905), mRNA
NM_014933	Homo sapiens yeast Sec31p homolog (KIAA0905), mRNA
NM_014706	Homo sapiens squamous cell carcinoma antigen recognised by T cells 3
	(SART3), mRNA
NM_014463	Homo sapiens Lsm3 protein (LSM3), mRNA
NM_014288	Homo sapiens integrin beta 3 binding protein (beta3-endonexin) (ITGB3BP), mRNA
NM_013443	Homo sapiens CMP-NeuAC:(beta)-N-acetylgalactosaminide (alpha)2,6-sialyltransferase member VI (VI), mRNA
NM 012404	Homo sapiens pp32 related 2 (PP32R2), mRNA
NM 012403	Homo sapiens pp32 related 1 (PP32R1), mRNA
NM 006710	Homo sapiens COP9 homolog (COP9), mRNA
NM 006117	Homo sapiens peroxisomal D3,D2-enoyl-CoA isomerase (PECI), mRNA
NM_005839	Homo sapiens serine/arginine repetitive matrix 1 (SRRM1), mRNA
NM_004264	Homo sapiens SRB7 suppressor of RNA polymerase B homolog (yeast) (SURB7), mRNA
NM 003714	Homo sapiens stanniocalcin 2 (STC2), mRNA
NM_003122	Homo sapiens serine protease inhibitor, Kazal type 1 (SPINK1), mRNA
NM_003690	Homo sapiens protein kinase, interferon-inducible double stranded RNA
	dependent activator (PRKRA), mRNA
NM_015526	Homo sapiens CLIP-170-related protein (CLIPR-59), mRNA
NM_033401	Homo sapiens cell recognition protein CASPR4 (CASPR4), mRNA
NM_023037	Homo sapiens hypothetical protein CG003 (13CDNA73), mRNA
NM_021817	Homo sapiens brain link protein-1 (BRAL1), mRNA
NM_016222	Homo sapiens DEAD-box protein abstrakt (ABS), mRNA
NM 003744	Homo sapiens numb homolog (Drosophila) (NUMB), mRNA
NM_032682	Homo sapiens forkhead box P1 (FOXP1), mRNA
NM_003681	Homo sapiens pyridoxal (pyridoxine, vitamin B6) kinase (PDXK), mRNA

NTM 001605	1 **
NM_001685	Homo sapiens ATP synthase, H+ transporting, mitochondrial F0 complex,
NM 017954	subunit F6 (ATP5J), mRNA
NM 015626	Homo sapiens hypothetical protein FLJ20761 (FLJ20761), mRNA
NM_130795	Homo sapiens SOCS box-containing WD protein SWiP-1 (WSB1), mRNA
NM_030877	Homo sapiens regulator of G-protein signalling 3 (RGS3), mRNA
NM_080830	Homo sapiens chromosome 20 open reading frame 33 (C20orf33), mRNA
NM_032329	Homo sapiens cystatin 11 (CST11), mRNA
NM_022917	Homo sapiens p28 ING5 (ING5), mRNA
NM_130787	Homo sapiens nucleolar RNA-associated protein (Nrap), mRNA
	Homo sapiens adaptor-related protein complex 2, alpha 1 subunit (AP2A1), mRNA
NM_024744	Homo sapiens (ALS2CR8), mRNA
NM 018984	Homo sapiens slingshot 1 (hSSH-1), mRNA
NM_106552	Homo sapiens hypothetical protein FLJ14249 similar to HS1 binding protein 3 (FLJ14249), transcript variant 2, mRNA
NM_022460	Homo sapiens hypothetical protein FLJ14249 similar to HS1 binding protein 3
	(FLJ14249), transcript variant 1, mRNA
NM_130446	Homo sapiens kelch-like protein KLHL6 (KLHL6), mRNA
NM_020314	Homo sapiens esophageal cancer associated protein (MGC16824) mRNA
NM_130395	Homo sapiens Werner helicase interacting protein (WHIP), transcript variant 2
	IIIRNA
NM_020135	Homo sapiens Werner helicase interacting protein (WHIP), transcript variant 1,
	IIIRIVA
NM_130388	Homo sapiens ankyrin repeat and SOCS box-containing 12 (ASB12), mRNA
NM_130387	Homo sapiens ankyrin repeat and SOCS box-containing 14 (ASB14), mRNA
NM_007191	Homo sapiens WN1 inhibitory factor 1 (WIF1) mRNA
NM_052950	Homo sapiens WD40- and FYVE-domain containing protein 2 (WDF2), mPNA
NM_025042	mRNA withams-Beuren syndrome chromosome region 23 (WBSCR23),
NM_080706	Homo sapiens transient receptor potential cation channel, subfamily V, member
	1 (TKP v I), transcript variant 3, mRNA
NM_080705	Homo sapiens transient receptor potential cation channel subfamily V member
37.000	1 (TRPVI), transcript variant 4, mRNA
NM_080704	Homo sapiens transient receptor potential cation channel, subfamily V, member
) To 6 010707	1 (TRP VI), transcript variant I, mRNA
NM_018727	Homo sapiens transient receptor potential cation channel, subfamily V, member
ND 4 000070	1 (TRPVI), transcript variant 2, mRNA
NM_080879	Homo sapiens SOCS box containing protein RAR2A (RAR2A), mRNA
NM 080871	Homo sapiens ankyrin repeat and SOCS box-containing 10 (ASB10), mRNA
NM_080870	riomo sapiens DPCR1 protein (DPCR1), mRNA
NM_080834 NM_080829	Homo sapiens chromosome 20 open reading frame 152 (C20orf152), mRNA
NM 080828	Homo sapiens chromosome 20 open reading frame 175 (C20orf175) mRNA
NM 080819	Homo sapiens chromosome 20 open reading frame 173 (C20orf173), mRNA
NM_080752	Homo sapiens G protein-coupled receptor 78 (GPR78), mRNA
NM_080749	Homo sapiens chromosome 20 open reading frame 164 (C20orf164), mRNA
NM 080745	Homo sapiens chromosome 20 open reading frame 163 (C20orf163), mRNA
NM_080738	Homo sapiens ring finger protein 36 (RNF36), mRNA
NM_014970	Homo sapiens EDAR-associated death domain (EDARADD), mRNA
NM_021058	Homo sapiens kinesin-associated protein 3 (KIFAP3), mRNA
NM_021058	Homo sapiens H2B histone family, member R (H2BFR), mRNA
NM_080491	Homo sapiens GPP2
1.1.1_000-191	Homo sapiens GRB2-associated binding protein 2 (GAB2), transcript variant 1,

	mRNA
NM_012296	Homo sapiens GRB2-associated binding protein 2 (GAB2), transcript variant 2,
14141_012250	mRNA
NM 007247	Homo sapiens AP1 gamma subunit binding protein 1 (AP1GBP1), transcript
14141_007247	variant 1, mRNA
NTM 000551	Homo sapiens AP1 gamma subunit binding protein 1 (AP1GBP1), transcript
NM_080551	variant 3, mRNA
NT 000550	Homo sapiens AP1 gamma subunit binding protein 1 (AP1GBP1), transcript
NM_080550	
	variant 2, mRNA Homo sapiens ribosomal protein L21 (RPL21), mRNA
NM_000982	Homo sapiens ribosomar protein L21 (Kl L21), indext. Homo sapiens serine/threonine-protein kinase PRP4 homolog (PRP4), mRNA
NM_003913	Homo sapiens senne/inreonine-protein kinase I KI 4 homolog (I KI 4), mkd VI
NM_002475	Homo sapiens myosin light chain 1 slow a (MLC1SA), mRNA
NM_002729	Homo sapiens hematopoietically expressed homeobox (HHEX), mRNA
NM_005893	Homo sapiens calicin (CCIN), mRNA
NM_017593	Homo sapiens homolog of mouse BMP-2 inducible kinase (BIKE), mRNA
NM_032027	Homo sapiens beta-amyloid binding protein precursor (BBP), mRNA
NM_004051	Homo sapiens 3-hydroxybutyrate dehydrogenase (heart, mitochondrial) (BDH),
	nuclear gene encoding mitochondrial protein, mRNA
NM 006576	Homo sapiens advillin (AVIL), mRNA
NM 013375	Homo sapiens TATA-binding protein-binding protein (ABT1), mRNA
NM 058219	Homo sapiens homolog of yeast mRNA transport regulator 3 (MTR3), mRNA
NM 058237	Homo sapiens HEAT-like repeat-containing protein (KIAA1622), transcript
· –	variant 1, mRNA
NM_020958	Homo sapiens HEAT-like repeat-containing protein (KIAA1622), transcript
* .2 <u> </u>	variant 2, mRNA
NM 004702	Homo sapiens cyclin E2 (CCNE2), transcript variant 3, mRNA
NM 057749	Homo sapiens cyclin E2 (CCNE2), transcript variant 1, mRNA
NM 057735	Homo sapiens cyclin E2 (CCNE2), transcript variant 2, mRNA
NM 002013	Homo sapiens FK506 binding protein 3 (25kD) (FKBP3), mRNA
NM 004724	Homo sapiens ZW10 homolog, centromere/kinetochore protein (Drosophila)
14112_00 1721	(ZW10), mRNA
NM 057159	Homo sapiens endothelial differentiation, lysophosphatidic acid G-protein-
14141_057157	coupled receptor, 2 (EDG2), transcript variant 2, mRNA
NM_001401	Homo sapiens endothelial differentiation, lysophosphatidic acid G-protein-
14147_001401	coupled receptor, 2 (EDG2), transcript variant 1, mRNA
NM_015084	Homo sapiens mitochondrial ribosomal protein S27 (MRPS27), nuclear gene
14147_012004	encoding mitochondrial protein, mRNA
NM_033281	Homo sapiens mitochondrial ribosomal protein S36 (MRPS36), nuclear gene
14141_033261	encoding mitochondrial protein, mRNA
NM 005830	Homo sapiens mitochondrial ribosomal protein S31 (MRPS31), nuclear gene
14141_002920	encoding mitochondrial protein, mRNA
NM 012062	Homo sapiens dynamin 1-like (DNM1L), transcript variant 1, mRNA
NM 005648	Homo sapiens transcription elongation factor B (SIII), polypeptide 1 (15kD,
MM_003046	elongin C) (TCEB1), mRNA
NR 6 007070	Homo sapiens FKBP-associated protein (FAP48), transcript variant 2, mRNA
NM_007070	Homo sapiens FKBP-associated protein (FAP48), transcript variant 1, mRNA
NM_053274	HOMO Sapiens PND Amount - t mentain language anti-latin gubunit interacting
NM_054113	Homo sapiens DNA-dependent protein kinase catalytic subunit-interacting
	protein 3 (KIP3), mRNA
NM 003726	Homo sapiens src family associated phosphoprotein 1 (SCAP1), mRNA
NM_012308	Homo sapiens F-box and leucine-rich repeat protein 11 (FBXL11), mRNA
NM_030913	Homo sapiens sema domain, transmembrane domain (TM), and cytoplasmic
	domain, (semaphorin) 6C (SEMA6C), mRNA

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NM 021163	Homo sapiens RB-associated KRAB repressor (RBAK), mRNA
NM_033632	Homo sapiens F-box and WD-40 domain protein 7 (archipelago homolog,
	Drosophila) (FBXW7), transcript variant 1, mRNA
NM_018315	Homo sapiens F-box and WD-40 domain protein 7 (archipelago homolog,
-	Drosophila) (FBXW7), transcript variant 2, mRNA
NM_012168	Homo sapiens F-box only protein 2 (FBXO2), mRNA
NM_033332	Homo sapiens CDC14 cell division cycle 14 homolog B (S. cerevisiae)
	(CDC14B), transcript variant 3, mRNA
NM_033331	Homo sapiens CDC14 cell division cycle 14 homolog B (S. cerevisiae)
	(CDC14B), transcript variant 2, mRNA
NM_003671	Homo sapiens CDC14 cell division cycle 14 homolog B (S. cerevisiae)
	(CDC14B), transcript variant 1, mRNA
NM_033307	Homo sapiens caspase 4, apoptosis-related cysteine protease (CASP4), transcript
	variant delta, mRNA
NM_033306	Homo sapiens caspase 4, apoptosis-related cysteine protease (CASP4), transcript
	variant gamma, mRNA
NM_001225	Homo sapiens caspase 4, apoptosis-related cysteine protease (CASP4), transcript
	variant alpha, mRNA
NM_002948	Homo sapiens ribosomal protein L15 (RPL15), mRNA
NM_033228	Homo sapiens ADP-ribosylation factor domain protein 1, 64kD (ARFD1),
	transcript variant gamma, mRNA
NM_033227	Homo sapiens ADP-ribosylation factor domain protein 1, 64kD (ARFD1),
	transcript variant beta, mRNA
NM_001656	Homo sapiens ADP-ribosylation factor domain protein 1, 64kD (ARFD1),
	transcript variant alpha, mRNA
NM_021203	Homo sapiens APMCF1 protein (APMCF1), mRNA
NM_012095	Homo sapiens adaptor-related protein complex 3, mu 1 subunit (AP3M1),
	mRNA
NM_001025	Homo sapiens ribosomal protein S23 (RPS23), mRNA
NM_032989	Homo sapiens BCL2-antagonist of cell death (BAD), transcript variant 2, mRNA
NM_004322	Homo sapiens BCL2-antagonist of cell death (BAD), transcript variant 1, mRNA
NM_014326	Homo sapiens death-associated protein kinase 2 (DAPK2), mRNA
NM_012430	Homo sapiens sec22 homolog (SEC22A), mRNA
NM_031216	Homo sapiens sec13-like protein (SEC13L), mRNA
NM_002927	Homo sapiens regulator of G-protein signalling 13 (RGS13), mRNA
NM_031274	Homo sapiens testis expressed sequence 13A (TEX13A), mRNA
NM_001730	Homo sapiens Kruppel-like factor 5 (intestinal) (KLF5), mRNA
NM_032674	Homo sapiens leucine rich repeat (in FLII) interacting protein 1 (LRRFIP1),
	mRNA
NM_031361	Homo sapiens collagen, type IV, alpha 3 (Goodpasture antigen) binding protein
	(COL4A3BP), transcript variant 2, mRNA
NM_031266	Homo sapiens heterogeneous nuclear ribonucleoprotein A/B (HNRPAB),
-	transcript variant 1, mRNA
NM_004499	Homo sapiens heterogeneous nuclear ribonucleoprotein A/B (HNRPAB),
·	transcript variant 2, mRNA
NM_004990	Homo sapiens methionine-tRNA synthetase (MARS), mRNA
NM_031244	Homo sapiens sirtuin silent mating type information regulation 2 homolog 5 (S.
_	cerevisiae) (SIRT5), transcript variant 2, mRNA
NM_012241	Homo sapiens sirtuin silent mating type information regulation 2 homolog 5 (S.
-	cerevisiae) (SIRT5), transcript variant 1, mRNA
NM_006845	Homo sapiens kinesin-like 6 (mitotic centromere-associated kinesin) (KNSL6),
_	mRNA

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Homo sapiens lecuine-rich acidic protein-like protein (LANP-L), mRNA
Homo sapiens L-kynurenine/alpha-aminoadipate aminotransferase (KATII),
mRNA
Homo sapiens hypothetical protein FLJ20297 (FLJ20297), mRNA
Homo sapiens cytochrome P450, subfamily IVA, polypeptide 11 (CYP4A11), mRNA
Homo sapiens glucocorticoid modulatory element binding protein 1 (GMEB1), transcript variant 1, mRNA
Homo sapiens glucocorticoid modulatory element binding protein 1 (GMEB1), transcript variant 2, mRNA
Homo sapiens TAF7-like RNA polymerase II, TATA box binding protein (TBP)-associated factor, 50 kD (TAF7L), mRNA
Homo sapiens ARP1 actin-related protein 1 homolog A, centractin alpha (yeast) (ACTR1A), mRNA
Homo sapiens VLCS-H1 protein (VLCS-H1), mRNA
Homo sapiens integrin cytoplasmic domain-associated protein 1 (ICAP-1A), transcript variant 2, mRNA
Homo sapiens endothelial cell-specific molecule 1 (ESM1), mRNA
Homo sapiens chromosome 12 open reading frame 8 (C12orf8), mRNA
Homo sapiens C-terminal binding protein 2 (CTBP2), transcript variant 2, mRNA
Homo sapiens E2F transcription factor 5, p130-binding (E2F5), mRNA
Homo sapiens epididymal sperm binding protein 1 (ELSPBP1), mRNA
Homo sapiens beta-1,3-glucuronyltransferase 3 (glucuronosyltransferase I) (B3GAT3), mRNA
Homo sapiens oculomedin (OCLM), mRNA
Homo sapiens growth differentiation factor 10 (GDF10), mRNA
Homo sapiens RNA helicase-related protein (RNAHP), mRNA
Homo sapiens regulator of G-protein signalling 4 (RGS4), mRNA
Homo sapiens IK cytokine, down-regulator of HLA II (IK), mRNA
Homo sapiens splicing factor 3b, subunit 3, 130kD (SF3B3), mRNA
Homo sapiens hypothetical protein FLJ10637 (FLJ10637), mRNA
Homo sapiens adenylyl cyclase-associated protein (CAP), mRNA
Homo sapiens regulator of G-protein signalling 3 (RGS3), mRNA
Homo sapiens solute carrier family 15 (H+/peptide transporter), member 2 (SLC15A2), mRNA
Homo sapiens HBV pX associated protein-8 (LOC51773), mRNA
Homo sapiens solute carrier family 1 (glutamate transporter), member 7 (SLC1A7), mRNA
Homo sapiens hypothetical protein LOC57333 (LOC57333), mRNA
Homo sapiens lymphocyte activation-associated protein (LOC51088), mRNA
Homo sapiens PAN2 protein (PAN2), mRNA
Homo sapiens HT021 (HT021), mRNA
Homo sapiens Cyt19 protein (Cyt19), mRNA
Homo sapiens HT017 protein (HT017), mRNA
Homo sapiens uncharacterized gastric protein ZA52P (LOC57399), mRNA
Homo sapiens eukaryotic translation initiation factor 4 gamma, 3 (EIF4G3), mRNA
IIIXIVX
Homo sapiens CHMP1.5 protein (CHMP1.5), mRNA

77.6 00000	
NM_020387	Homo sapiens CATX-8 protein (CATX-8), mRNA
NM_020371	Homo sapiens cell death regulator aven (LOC57099), mRNA
NM_020362	Homo sapiens HT014 (HT014), mRNA
NM_020307	Homo sapiens cyclin L ania-6a (LOC57018), mRNA
NM_007187	Homo sapiens WW domain binding protein 4 (formin binding protein 21) (WBP4), mRNA
NM_005644	Homo sapiens TAF12 RNA polymerase II, TATA box binding protein (TBP)-associated factor, 20 kD (TAF12), mRNA
NM 020150	Homo sapiens SAR1 protein (SAR1), mRNA
NM 020167	Homo sapiens neuromedin U receptor 2 (NMU2R), mRNA
NM 020233	Homo sapiens x 006 protein (MDS006), mRNA
NM_020232	Homo sapiens x 003 protein (MDS003), mRNA Homo sapiens x 003 protein (MDS003), mRNA
NM 020247	Homo sapiens hypothetical protein, clone
1117_020247	Telethon(Italy B41) Strait02270 FL142 (LOC56997), mRNA
NM_020213	Homo sapiens hypothetical protein from EUROIMAGE 1977056 (LOC56965),
	mRNA
NM_020153	Homo sapiens hypothetical protein (LOC56912), mRNA
NM_020149	Homo sapiens Meis1, myeloid ecotropic viral integration site 1 homolog 2 (mouse) (MEIS2), mRNA
NM_020120	Homo sapiens UDP-glucose ceramide glucosyltransferase-like 1 (UGCGL1), mRNA
NM_020190	Homo sapiens HNOEL-iso protein (HNOEL-iso), mRNA
NM_020242	Homo sapiens kinesin-like 7 (KNSL7), mRNA
NM_020194	Homo sapiens GL004 protein (GL004), mRNA
NM_020193	Homo sapiens GL002 protein (GL002), mRNA
NM_020189	Homo sapiens DC6 protein (DC6), mRNA
NM_020188	Homo sapiens DC13 protein (DC13), mRNA
NM_020134	Homo sapiens collapsin response mediator protein-5; CRMP3-associated molecule (CRMP5), mRNA
NM_019893	Homo sapiens mitochondrial ceramidase (ASAH2), mRNA
NM_019846	Homo sapiens CC chemokine CCL28 (SCYA28), mRNA
NM_019852	Homo sapiens putative methyltransferase (M6A), mRNA
NM_013338	Homo sapiens Alg5, S. cerevisiae, homolog of (ALG5), mRNA
NM 013341	Homo sapiens hypothetical protein (PTD004), mRNA
NM_013318	Homo sapiens hypothetical protein (LQFBS-1), mRNA
NM_013302	Homo sapiens elongation factor-2 kinase (HSU93850), mRNA
NM_013299	Homo sapiens protein predicted by clone 23627 (HSU79266), mRNA
NM_013347	Homo sapiens replication protein A complex 34 kd subunit homolog Rpa4 (HSU24186), mRNA
NM 019011	Homo sapiens TRIAD3 protein (TRIAD3), mRNA
NM_018965	Homo sapiens triggering receptor expressed on myeloid cells 2 (TREM2).
NM 019043	mRNA
NM_019006	Homo sapiens similar to proline-rich protein 48 (LOC54518), mRNA
NM 019101	Homo sapiens protein associated with PRK1 (AWP1), mRNA
NM 019049	Homo sapiens apolipoprotein M (G3A), mRNA
NM_018992	Homo sapiens hypothetical protein (FLJ20054), mRNA
NM_019033	Homo sapiens hypothetical protein (FLJ20040), mRNA
NM_019045	Homo sapiens hypothetical protein (FLJ11235), mRNA
NM_019079	Homo sapiens similar to rab11-binding protein (FLJ11116), mRNA
NM_019079	Homo sapiens hypothetical protein (FLJ10884), mRNA
NM_014298	Homo sapiens hypothetical protein (FLJ10007), mRNA
14141 014298	Homo sapiens quinolinate phosphoribosyltransferase (nicotinate-nucleotide

	pyrophosphorylase (carboxylating)) (QPRT), mRNA
NM 012413	Homo sapiens glutaminyl-peptide cyclotransferase (glutaminyl cyclase) (QPCT),
NM_012413	mRNA
NM 018836	Homo sapiens hypothetical protein (MOT8), mRNA
NM 018643	Homo sapiens triggering receptor expressed on myeloid cells 1 (TREM1),
	mRNA
NM_018647	Homo sapiens tumor necrosis factor receptor superfamily, member 19 (TNFRSF19), mRNA
NM 018664	Homo sapiens Jun dimerization protein p21SNFT (SNFT), mRNA
NM 018540	Homo sapiens hypothetical protein PRO2831 (PRO2831), mRNA
NM 018630	Homo sapiens hypothetical protein PRO2577 (PRO2577), mRNA
NM 018527	Homo sapiens hypothetical protein PRO2435 (PRO2435), mRNA
NM 018625	Homo sapiens hypothetical protein PRO2289 (PRO2289), mRNA
NM 018515	Homo sapiens hypothetical protein PRO2176 (PRO2176), mRNA
NM 018615	Homo sapiens hypothetical protein PRO2032 (PRO2032), mRNA
NM 018614	Homo sapiens hypothetical protein PRO2012 (PRO2012), mRNA
NM 018608	Homo sapiens hypothetical protein PRO1905 (PRO1905), mRNA
NM 018509	Homo sapiens hypothetical protein PRO1855 (PRO1855), mRNA
NM 018505	Homo sapiens hypothetical protein PRO1728 (PRO1728), mRNA
NM 018444	Homo sapiens pyruvate dehydrogenase phosphatase (PDP), mRNA
NM 018442	Homo sapiens PC326 protein (PC326), mRNA
NM 018698	Homo sapiens hypothetical protein P15-2 (P15-2), mRNA
NM_018466	Homo sapiens uncharacterized hematopoietic stem/progenitor cells protein
-	MDS031 (MDS031), mRNA
NM_018465	Homo sapiens uncharacterized hematopoietic stem/progenitor cells protein MDS030 (MDS030), mRNA
NM_018463	Homo sapiens uncharacterized hematopoietic stem/progenitor cells protein MDS028 (MDS028), mRNA
NM 018650	Homo sapiens MAP/microtubule affinity-regulating kinase 1 (MARK1), mRNA
NM 018678	Homo sapiens lipopolysaccharide specific response-68 protein (LSR68), mRNA
NM 018695	Homo sapiens erbb2 interacting protein (ERBB2IP), mRNA
NM_018683	Homo sapiens zinc finger protein 313 (ZNF313), mRNA
NM_018660	Homo sapiens papillomavirus regulatory factor PRF-1 (LOC55893), mRNA
NM_018484	Homo sapiens solute carrier family 22 (organic anion/cation transporter), member 11 (SLC22A11), mRNA
NM 018445	Homo sapiens AD-015 protein (LOC55829), mRNA
NM 017571	Homo sapiens hypothetical protein (LOC55580), mRNA
NM_017542	Homo sapiens KIAA1513 protein (KIAA1513), mRNA
NM_018473	Homo sapiens uncharacterized hypothalamus protein HT012 (HT012), mRNA
NM_018480	Homo sapiens uncharacterized hypothalamus protein HT007 (HT007), mRNA
NM_017583	Homo sapiens DIPB protein (HSA249128), mRNA
NM 017567	Homo sapiens N-acetylglucosamine kinase (NAGK), mRNA
NM_018487	Homo sapiens hepatocellular carcinoma-associated antigen 112 (HCA112), mRNA
NM 017548	Homo sapiens hypothetical protein (H41), mRNA
NM 017547	Homo sapiens hypothetical protein (H17), mRNA
NM 017966	Homo sapiens hypothetical protein (H17), linkina Homo sapiens hypothetical protein FLJ20847 (FLJ20847), mRNA
NM 017955	Homo sapiens hypothetical protein FLJ20847 (FLJ20847), mRNA
NM 017948	Homo sapiens hypothetical protein FLJ20764 (FLJ20764), mRNA Homo sapiens hypothetical protein FLJ20736 (FLJ20736), mRNA
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NM_017886	Homo sapiens hypothetical protein FLJ20574 (FLJ20574), mRNA
NM_017880	Homo sapiens hypothetical protein FLJ20558 (FLJ20558), mRNA
NM_017878	Homo sapiens HRAS-like suppressor 2 (HRASLS2), mRNA
NM_017877	Homo sapiens hypothetical protein FLJ20555 (FLJ20555), mRNA
NM_017875	Homo sapiens hypothetical protein FLJ20551 (FLJ20551), mRNA
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NM 017867	Homo sapiens hypothetical protein FLJ20534 (FLJ20534), mRNA
NM_017864	Homo sapiens hypothetical protein FLJ20530 (FLJ20530), mRNA
NM_017857	Homo sapiens slingshot 3 (SSH-3), mRNA
NM_017852	Homo sapiens NALP2 protein (NALP2), mRNA
NM 017850	Homo sapiens hypothetical protein FLJ20508 (FLJ20508), mRNA
NM 017846	Homo sapiens tRNA selenocysteine associated protein (SECP43), mRNA
NM 017841	Homo sapiens hypothetical protein FLJ20487 (FLJ20487), mRNA
NM 017839	Homo sapiens hypothetical protein FLJ20481 (FLJ20481), mRNA
NM 017837	Homo sapiens hypothetical protein FLJ20477 (FLJ20477), mRNA
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NM 017802	Homo sapiens hypothetical protein FLJ20397 (FLJ20397), mRNA
NM 017792	Homo sapiens hypothetical protein FLJ20373 (FLJ20373), mRNA
NM 017790	Homo sapiens regulator of G-protein signalling 3 (RGS3), mRNA
NM_017786	Homo sapiens hypothetical protein FLJ20366 (FLJ20366), mRNA
NM_017785	Homo sapiens hypothetical protein FLJ20364 (FLJ20364), mRNA
NM_017775	Homo sapiens hypothetical protein FLJ20343 (FLJ20343), mRNA
NM_017774	Homo sapiens hypothetical protein FLJ20342 (FLJ20342), mRNA
NM_017772	Homo sapiens hypothetical protein FLJ20337 (FLJ20337), mRNA
NM_017770	Homo sapiens elongation of very long chain fatty acids (FEN1/Elo2, SUR4/Elo3,
	yeast)-like 2 (ELOVL2), mRNA
NM_017762	Homo sapiens hypothetical protein FLJ20313 (FLJ20313), mRNA
NM_017759	Homo sapiens hypothetical protein FLJ20309 (FLJ20309), mRNA
NM_017756	Homo sapiens hypothetical protein FLJ20306 (FLJ20306), mRNA
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NM_017740	Homo sapiens hypothetical protein FLJ20279 (FLJ20279), mRNA
NM_017738	Homo sapiens hypothetical protein FLJ20276 (FLJ20276), mRNA

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NM_017735	Homo sapiens hypothetical protein FLJ20272 (FLJ20272), mRNA
NM_017719	Homo sapiens hypothetical protein FLJ20224 (FLJ20224), mRNA
NM_017718	Homo sapiens hypothetical protein FLJ20220 (FLJ20220), mRNA
NM_017716	Homo sapiens membrane-spanning 4-domains, subfamily A, member 12 4-
ND (017711	domains, subfamily A, member 7 (MS4A12), mRNA
NM_017711	Homo sapiens hypothetical protein FLJ20207 (FLJ20207), mRNA
NM_017709	Homo sapiens hypothetical protein FLJ20202 (FLJ20202), mRNA
NM_017704	Homo sapiens hypothetical protein FLJ20189 (FLJ20189), mRNA
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NM 017697	Homo sapiens hypothetical protein FLJ20171 (FLJ20171), mRNA
NM_017687	Homo sapiens hypothetical protein FLJ20147 (FLJ20147), mRNA
NM_017686	Homo sapiens ganglioside induced differentiation associated protein 2 (GDAP2), mRNA
NM_017678	Homo sapiens hypothetical protein FLJ20127 (FLJ20127), mRNA
NM_017677	Homo sapiens hypothetical protein FLJ20126 (FLJ20126), mRNA
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NM_017645	Homo sapiens hypothetical protein FLJ20060 (FLJ20060), mRNA
NM_017640	Homo sapiens hypothetical protein FLJ20048 (FLJ20048), mRNA
NM_017637	Homo sapiens hypothetical protein FLJ20043 (FLJ20043), mRNA
NM_017636	Homo sapiens transient receptor potential cation channel, subfamily M, member 4 (TRPM4), mRNA
NM 017634	Homo sapiens hypothetical protein FLJ20038 (FLJ20038), mRNA
NM 017629	Homo sapiens hypothetical protein FLJ20033 (FLJ20033), mRNA
NM_017622	Homo sapiens hypothetical protein FLJ20014 (FLJ20014), mRNA
NM_017620	Homo sapiens hypothetical protein FLJ20011 (FLJ20011), mRNA
NM_018396	Homo sapiens putative methyltransferase (METL), mRNA
NM_018381	Homo sapiens hypothetical protein FLJ11286 (FLJ11286), mRNA
NM_018371	Homo sapiens hypothetical protein FLJ11264 (FLJ11264), mRNA
NM_018368	Homo sapiens hypothetical protein FLJ11240 (FLJ11240), mRNA
NM_018367	Homo sapiens phytoceramidase, alkaline (PHCA), mRNA
NM_018364	Homo sapiens hypothetical protein FLJ11220 (FLJ11220), mRNA
NM_018363	Homo sapiens hypothetical protein FLJ11218 (FLJ11218), mRNA
NM_018361	Homo sapiens hypothetical protein FLJ11210 (FLJ11210), mRNA
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NM_018333	Homo sapiens hypothetical protein FLJ20666 (FLJ20666), mRNA
NM_018332	Homo sapiens hypothetical protein FLJ11126 (FLJ11126), mRNA
NM_018330	Homo sapiens KIAA1598 protein (KIAA1598), mRNA
NM_018322	Homo sapiens hypothetical protein FLJ11101 (FLJ11101), mRNA
NM_018318	Homo sapiens hypothetical protein FLJ11088 (FLJ11088), mRNA
NM_018310	Homo sapiens BRF2, subunit of RNA polymerase III transcription initiation
	factor, BRF1-like (BRF2), mRNA

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NM_018489	Homo sapiens chromosome 6 open reading frame 35 (C6orf35), mRNA
NM_004227	Homo sapiens hypothetical protein ASH1 (ASH1), mRNA
<u>-</u> 007227	Homo sapiens pleckstrin homology, Sec7 and coiled/coil domains 3 (PSCD3), mRNA
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NM_017431	Homo sapiens Nedd-4-like ubiquitin-protein ligase (WWP2), mRNA
	Homo sapiens protein kinase, AMP-activated, gamma 3 non-catalytic subunit

	(PRKAG3), mRNA
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NM_017421	Homo sapiens methyltransferase COQ3 (COQ3), mRNA Homo sapiens KDEL (Lys-Asp-Glu-Leu) endoplasmic reticulum protein
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NM_016524	Homo sapiens B/K protein (LOC51760), mRNA
NM_016507	Homo sapiens CDC2-related protein kinase 7 (CrkRS), mRNA
NM_016446	Homo sapiens NAG-5 protein (LOC51754), mRNA
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NM 016261	Homo sapiens delta-tubulin (LOC51174), mRNA
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NM 016208	Homo sapiens VPS28 protein (LOC51160), mRNA
NM 016206	Homo sapiens colon carcinoma related protein (LOC51159), mRNA
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NM_016185	Homo sapiens hematological and neurological expressed 1 (HN1), mRNA
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NM 014778	Homo sapiens KIAA0410 gene product (KIAA0410), mRNA
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NM 014720	Homo sapiens Ste20-related serine/threonine kinase (SLK), mRNA
NM 014761	Homo sapiens KIAA0174 gene product (KIAA0174), mRNA
NM 014730	Homo sapiens KIAA0152 gene product (KIAA0152), mRNA
NM 014661	Homo sapiens KIAA0140 gene product (KIAA0140), mRNA
NM 014777	Homo sapiens KIAA0133 gene product (KIAA0133), mRNA
NM_014815	Homo sapiens KIAA0130 gene product (KIAA0130), mRNA
NM_014755	Homo sapiens transcriptional regulator interacting with the PHS-bromodomain 2
	(TRIP-Br2), mRNA
NM 014628	Homo sapiens gene predicted from cDNA with a complete coding sequence
	(KIAA0110), mRNA
NM 014814	Homo sapiens KIAA0107 gene product (KIAA0107), mRNA
NM 014752	Homo sapiens KIAA0102 gene product (KIAA0102), mRNA
NM 014780	Homo sapiens KIAA0076 gene product (KIAA0076), mRNA
NM 014882	Homo sapiens KIAA0053 gene product (KIAA0053), mRNA
NM 014750	Homo sapiens KIAA0008 gene product (KIAA0008), mRNA
NM 015684	Homo sapiens mitochondrial ATP synthase regulatory component factor B
	(ATPW), mRNA
NM 014186	Homo sapiens HSPC166 protein (HSPC166), mRNA
NM_014184	Homo sapiens HSPC163 protein (HSPC163), mRNA
NM 014181	Homo sapiens HSPC159 protein (HSPC159), mRNA
NM 014179	Homo sapiens HSPC157 protein (HSPC157), mRNA
NM 014166	Homo sapiens HSPC126 protein (HSPC126), mRNA
NM 014155	Homo sapiens HSPC063 protein (HSPC063), mRNA
NM 014038	Homo sapiens HSPC028 protein (HSPC028), mRNA
NM 014017	Homo sapiens HSPC003 protein (HSPC003), mRNA
NM 014053	Homo sapiens FLVCR protein (FLVCR), mRNA
NM 015400	Homo sapiens DKFZP586N0721 protein (DKFZP586N0721), mRNA
NM 015583	Homo sapiens DKFZP586M0622 protein (DKFZP586M0622), mRNA
NM 015485	Homo sapiens DKFZP566K023 protein (DKFZP566K023), mRNA
NM 014043	Homo sapiens DKFZP564O123 protein (DKFZP564O123), mRNA
NM 015387	Homo sapiens preimplantation protein 3 (PREI3), mRNA
NM_014056	Homo sapiens DKFZP564K247 protein (DKFZP564K247), mRNA
NM 015623	Homo sapiens putative ankyrin-repeat containing protein (DKFZP564D166),
	mRNA
NM 015582	Homo sapiens DKFZP564B147 protein (DKFZP564B147), mRNA
NM 015610	Homo sapiens DKFZP434J154 protein (DKFZP434J154), mRNA
NM 015590	Homo sapiens DKFZP434F1735 protein (DKFZP434F1735), mRNA
NM 015644	Homo sapiens DKFZP434B103 protein (DKFZP434B103), mRNA
NM 015396	Homo sapiens DKFZP434A043 protein (DKFZP434A043), mRNA
NM 014058	Homo sapiens DESC1 protein (DESC1), mRNA
NM_015680	Homo sapiens hypothetical protein (CGI-57), mRNA
NM 015379	Homo sapiens brain protein I3 (BRI3), mRNA
NM_014580	Homo sapiens solute carrier family 2, (facilitated glucose transporter) member 8
1111_014500	(SLC2A8), mRNA
NM 014280	Homo sapiens DnaJ (Hsp40) homolog, subfamily C, member 8 (DNAJC8),

	mRNA
NM 014313	Homo sapiens small membrane protein 1 (SMP1), mRNA
NM 014229	Homo sapiens solute carrier family 6 (neurotransmitter transporter, GABA),
1111_014225	member 11 (SLC6A11), mRNA
NM 014575	Homo sapiens schwannomin interacting protein 1 (SCHIP1), mRNA
NM 014402	Homo sapiens low molecular mass ubiquinone-binding protein (9.5kD) (QP-C),
_	mRNA
NM_014394	Homo sapiens growth hormone inducible transmembrane protein (GHITM), mRNA
NM_014225	Homo sapiens protein phosphatase 2 (formerly 2A), regulatory subunit A (PR 65), alpha isoform (PPP2R1A), mRNA
NM_014497	Homo sapiens nuclear protein (NP220), mRNA
NM_014399	Homo sapiens tetraspan NET-6 protein (NET-6), mRNA
NM_014889	Homo sapiens metalloprotease 1 (pitrilysin family) (MP1), mRNA
NM 014484	Homo sapiens molybdenum cofactor synthesis 3 (MOCS3), mRNA
NM 014447	Homo sapiens arfaptin 1 (HSU52521), mRNA
NM 014350	Homo sapiens TNF-induced protein (GG2-1), mRNA
NM_014478	Homo sapiens calcitonin gene-related peptide-receptor component protein (CGRP-RCP), mRNA
NM 014482	Homo sapiens bone morphogenetic protein 10 (BMP10), mRNA
NM_014474	Homo sapiens acid sphingomyelinase-like phosphodiesterase (ASML3B), mRNA
NM 014480	Homo sapiens zinc finger protein (AF020591), mRNA
NM_014576	Homo sapiens Apobec-1 complementation factor; APOBEC-1 stimulating protein (ACF), mRNA
NM 005884	Homo sapiens p21(CDKN1A)-activated kinase 4 (PAK4), mRNA
NM_013434	Homo sapiens calsenilin, presenilin binding protein, EF hand transcription factor (CSEN), mRNA
NM 012446	Homo sapiens single-stranded DNA binding protein 2 (SSBP2), mRNA
NM_013235	Homo sapiens putative ribonuclease III (RNASE3L), mRNA
NM 013349	Homo sapiens secreted protein of unknown function (SPUF), mRNA
NM_013323	Homo sapiens sorting nexin 11 (SNX11), mRNA
NM 013388	Homo sapiens prolactin regulatory element binding (PREB), mRNA
NM 013328	Homo sapiens pyrroline 5-carboxylate reductase isoform (P5CR2), mRNA
NM 013370	Homo sapiens pregnancy-induced growth inhibitor (OKL38), mRNA
NM 013277	Homo sapiens Rac GTPase activating protein 1 (RACGAP1), mRNA
NM 013285	Homo sapiens nucleolar GTPase (HUMAUANTIG), mRNA
NM 013320	Homo sapiens host cell factor 2 (HCF-2), mRNA
NM 013391	Homo sapiens dimethylglycine dehydrogenase precursor (DMGDH), mRNA
NM 013253	Homo sapiens dickkopf homolog 3 (Xenopus laevis) (DKK3), mRNA
NM_013339	Homo sapiens dolichyl-P-Glc:Man9GlcNAc2-PP-dolichylglucosyltransferase (ALG6), mRNA
NM 004120	Homo sapiens guanylate binding protein 2, interferon-inducible (GBP2), mRNA
NM 005690	Homo sapiens dynamin 1-like (DNM1L), transcript variant 3, mRNA
NM_012063	Homo sapiens dynamin 1-like (DNM1L), transcript variant 2, mRNA
NM 012470	Homo sapiens transportin-SR (TRN-SR), mRNA
NM 012252	Homo sapiens transcription factor EC (TFEC), mRNA
NM 012250	Homo sapiens related RAS viral (r-ras) oncogene homolog 2 (RRAS2), mRNA
NM_012249	Homo sapiens ras-like protein (TC10), mRNA
NM_012388	Homo sapiens pallidin homolog (mouse) (PLDN), mRNA
NM 012322	Homo sapiens U6 snRNA-associated Sm-like protein (LSM5), mRNA
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NM 012189	Homo sapiens fibrousheathin II (FSP-2), mRNA
NM 012081	Homo sapiens ELL-RELATED RNA POLYMERASE II, ELONGATION
14141_015001	FACTOR (ELL2), mRNA
NM 003996	Homo sapiens glutathione peroxidase 5 (epididymal androgen-related protein)
טבבנטס"זאזאז	(GPX5), transcript variant 2, mRNA
NM 005260	Homo sapiens growth differentiation factor 9 (GDF9), mRNA
NM 007352	Homo sapiens elastase 3B, pancreatic (ELA3B), mRNA
NM 006685	Homo sapiens proline rich 3 (PROL3), mRNA
NM_007357	Homo sapiens low density lipoprotein receptor defect C complementing (LDLC),
	mRNA
NM_004133	Homo sapiens hepatocyte nuclear factor 4, gamma (HNF4G), mRNA
NM_003144	Homo sapiens signal sequence receptor, alpha (translocon-associated protein alpha) (SSR1), mRNA
NM 007324	Homo sapiens MAD, mothers against decapentaplegic homolog (Drosophila)
	interacting protein, receptor activation anchor (MADHIP), transcript variant 1, mRNA
NM 007323	Homo sapiens MAD, mothers against decapentaplegic homolog (Drosophila)
	interacting protein, receptor activation anchor (MADHIP), transcript variant 2,
	mRNA
NM_005162	Homo sapiens angiotensin receptor-like 2 (AGTRL2), mRNA
NM_005501	Homo sapiens integrin, alpha 3 (antigen CD49C, alpha 3 subunit of VLA-3
	receptor) (ITGA3), transcript variant b, mRNA
NM_007144	Homo sapiens zinc finger protein 144 (Mel-18) (ZNF144), mRNA
NM_007286	Homo sapiens synaptopodin (KIAA1029), mRNA
NM_007199	Homo sapiens interleukin-1 receptor-associated kinase M (IRAK-M), mRNA
NM_007283	Homo sapiens monoglyceride lipase (MGLL), mRNA
NM_007241	Homo sapiens EAP30 subunit of ELL complex (EAP30), mRNA
NM_007212	Homo sapiens ring finger protein 2 (RNF2), mRNA
NM_007236	Homo sapiens calcium binding protein P22 (CHP), mRNA
NM_007063	Homo sapiens vascular Rab-GAP/TBC-containing (VRP), mRNA
NM_007027	Homo sapiens topoisomerase (DNA) II binding protein (TOPBP1), mRNA
NM_006938	Homo sapiens small nuclear ribonucleoprotein D1 polypeptide (16kD) (SNRPD1), mRNA
NM_006937	Homo sapiens SMT3 suppressor of mif two 3 homolog 2 (yeast) (SMT3H2),
	mRNA
NM_007029	Homo sapiens stathmin-like 2 (STMN2), mRNA
NM_007042	Homo sapiens ribonuclease P (14kD) (RPP14), mRNA
NM_006907	Homo sapiens pyrroline-5-carboxylate reductase 1 (PYCR1), nuclear gene
	encoding mitochondrial protein, mRNA
NM_007059	Homo sapiens kaptin (actin binding protein) (KPTN), mRNA
NM_007069	Homo sapiens HRAS-like suppressor 3 (HRASLS3), mRNA
NM_006895	Homo sapiens histamine N-methyltransferase (HNMT), mRNA
NM_007071	Homo sapiens HERV-H LTR-associating 3 (HHLA3), mRNA
NM_007067	Homo sapiens histone acetyltransferase (HBOA), mRNA
NM_007006	Homo sapiens cleavage and polyadenylation specific factor 5, 25 kD subunit
NM 007063	(CPSF5), mRNA Homo sapiens natural killer cell receptor, immunoglobulin superfamily member
NM_007053	(BY55), mRNA
NM_006754	Homo sapiens synaptophysin-like protein (SYPL), mRNA
NM_006802	Homo sapiens splicing factor 3a, subunit 3, 60kD (SF3A3), mRNA
NM_006842	Homo sapiens splicing factor 3b, subunit 2, 145kD (SF3B2), mRNA
NM_006834	Homo sapiens RAB32, member RAS oncogene family (RAB32), mRNA

NM 006875	Homo sapiens pim-2 oncogene (PIM2), mRNA
NM 006810	Homo sapiens for protein disulfide isomerase-related (PDIR), mRNA
NM 003609	Homo sapiens HIRA interacting protein 3 (HIRIP3), mRNA
NM 006820	Homo sapiens chromosome 1 open reading frame 29 (C1orf29), mRNA
NM 006848	Homo sapiens hepatitis delta antigen-interacting protein A (DIPA), mRNA
NM 006876	Homo sapiens UDP-GlcNAc:betaGal beta-1,3-N-acetylglucosaminyltransferase
_	6 (B3GNT6), mRNA
NM 006653	Homo sapiens suc1-associated neurotrophic factor target 2 (FGFR signalling
-	adaptor) (SNT-2), mRNA
NM 006638	Homo sapiens ribonuclease P, 40kD subunit (RPP40), mRNA
NM 004163	Homo sapiens RAB27B, member RAS oncogene family (RAB27B), mRNA
NM 006713	Homo sapiens activated RNA polymerase II transcription cofactor 4 (PC4),
_	mRNA .
NM 006601	Homo sapiens unactive progesterone receptor, 23 kD (P23), mRNA
NM 006675	Homo sapiens tetraspan transmembrane 4 super family (NET-5), mRNA
NM 006501	Homo sapiens myelin-associated oligodendrocyte basic protein (MOBP), mRNA
NM 006612	Homo sapiens kinesin family member 1C (KIF1C), mRNA
NM 006567	Homo sapiens phenylalanine-tRNA synthetase (FARS1), nuclear gene encoding
_	mitochondrial protein, mRNA
NM 006594	Homo sapiens adaptor-related protein complex 4, beta 1 subunit (AP4B1),
	mRNA
NM_006621	Homo sapiens S-adenosylhomocysteine hydrolase-like 1 (AHCYL1), mRNA
NM 006472	Homo sapiens thioredoxin interacting protein (TXNIP), mRNA
NM_006388	Homo sapiens HIV-1 Tat interactive protein, 60 kD (HTATIP), mRNA
NM 006281	Homo sapiens serine/threonine kinase 3 (STE20 homolog, yeast) (STK3),
-	mRNA
NM_006401	Homo sapiens acidic protein rich in leucines (SSP29), mRNA
NM_006425	Homo sapiens step II splicing factor SLU7 (SLU7), mRNA
NM_006359	Homo sapiens solute carrier family 9 (sodium/hydrogen exchanger), isoform 6
	(SLC9A6), mRNA
NM_006328	Homo sapiens RNA binding motif protein 14 (RBM14), mRNA
NM_006466	Homo sapiens polymerase (RNA) III (DNA directed) polypeptide F (39 kD)
	(POLR3F), mRNA
NM_006467	Homo sapiens polymerase (RNA) III (DNA directed) (32kD) (RPC32), mRNA
NM_006397	Homo sapiens ribonuclease HI, large subunit (RNASEHI), mRNA
NM_006443	Homo sapiens putative c-Myc-responsive (RCL), mRNA
NM_006390	Homo sapiens RAN binding protein 8 (RANBP8), mRNA
NM_006256	Homo sapiens protein kinase C-like 2 (PRKCL2), mRNA
NM_006254	Homo sapiens protein kinase C, delta (PRKCD), mRNA
NM_006229	Homo sapiens pancreatic lipase-related protein 1 (PNLIPRP1), mRNA
NM_006319	Homo sapiens CDP-diacylglycerolinositol 3-phosphatidyltransferase
	(phosphatidylinositol synthase) (CDIPT), mRNA
NM_006219	Homo sapiens phosphoinositide-3-kinase, catalytic, beta polypeptide (PIK3CB),
	mRNA
NM_006346	Homo sapiens progesterone-induced blocking factor 1 (PIBF1), mRNA
NM_006473	Homo sapiens TAF6-like RNA polymerase II, p300/CBP-associated factor
	(PCAF)-associated factor, 65 kD (TAF6L), mRNA
NM_006396	Homo sapiens Sjogren's syndrome/scleroderma autoantigen 1 (SSSCA1), mRNA
NM_006428	Homo sapiens melanoma-associated antigen recognised by cytotoxic T
	lymphocytes (MAAT1), mRNA
NM_006475	Homo sapiens osteoblast specific factor 2 (fasciclin I-like) (OSF-2), mRNA
NM 006392	Homo sapiens nucleolar protein 5A (56kD with KKE/D repeat) (NOL5A),

	mRNA
NM_006417	Homo sapiens interferon-induced, hepatitis C-associated microtubular aggregate
	protein (44kD) (MTAP44), mRNA
NM_006405	Homo sapiens transmembrane 9 superfamily member 1 (TM9SF1), mRNA
NM_006471	Homo sapiens myosin, light polypeptide, regulatory, non-sarcomeric (20kD) (MLCB), mRNA
NM 006152	Homo sapiens lymphoid-restricted membrane protein (LRMP), mRNA
NM 006460	Homo sapiens HMBA-inducible (HIS1), mRNA
NM_006365	Homo sapiens transcriptional activator of the c-fos promoter (CROC4), mRNA
NM_006135	Homo sapiens capping protein (actin filament) muscle Z-line, alpha 1 (CAPZA1), mRNA
NM 006086	Homo sapiens tubulin, beta, 4 (TUBB4), mRNA
NM 005761	Homo sapiens plexin C1 (PLXNC1), mRNA
NM 005724	Homo sapiens tetraspan 3 (TSPAN-3), mRNA
NM 005646	Homo sapiens TAR (HIV) RNA binding protein 1 (TARBP1), mRNA
NM 005819	Homo sapiens syntaxin 6 (STX6), mRNA
NM 005866	Homo sapiens sigma receptor (SR31747 binding protein 1) (SR-BP1), mRNA
NM 005842	Homo sapiens sprouty homolog 2 (Drosophila) (SPRY2), mRNA
NM 005626	Homo sapiens splicing factor, arginine/serine-rich 4 (SFRS4), mRNA
NM 005770	Homo sapiens small EDRK-rich factor 2 (SERF2), mRNA
NM 005805	Homo sapiens 26S proteasome-associated pad1 homolog (POH1), mRNA
NM 005746	Homo sapiens pre-B-cell colony-enhancing factor (PBEF), mRNA
NM_005869	Homo sapiens serologically defined colon cancer antigen 10 (SDCCAG10),
	mRNA
NM_005787	Homo sapiens Not56 (D. melanogaster)-like protein (NOT56L), mRNA
NM_005792	Homo sapiens M-phase phosphoprotein 6 (MPHOSPH6), mRNA
NM_005693	Homo sapiens nuclear receptor subfamily 1, group H, member 3 (NR1H3), mRNA
NM_005799	Homo sapiens PDZ domain protein (Drosophila inaD-like) (INADL), mRNA
NM_005713	Homo sapiens collagen, type IV, alpha 3 (Goodpasture antigen) binding protein (COL4A3BP), transcript variant 1, mRNA
NM 005878	Homo sapiens trinucleotide repeat containing 3 (TNRC3), mRNA
NM 005875	Homo sapiens translation factor sui1 homolog (GC20), mRNA
NM_005838	Homo sapiens glycine-N-acyltransferase (GLYAT), nuclear gene encoding mitochondrial protein, mRNA
NM_005754	Homo sapiens Ras-GTPase-activating protein SH3-domain-binding protein (G3BP), mRNA
NM_005764	Homo sapiens epithelial protein up-regulated in carcinoma, membrane associated protein 17 (DD96), mRNA
NM_005694	Homo sapiens COX17 homolog, cytochrome c oxidase assembly protein (yeast) (COX17), nuclear gene encoding mitochondrial protein, mRNA
NM_005506	Homo sapiens CD36 antigen (collagen type I receptor, thrombospondin receptor)-like 2 (lysosomal integral membrane protein II) (CD36L2), mRNA
NM_005881	Homo sapiens branched chain alpha-ketoacid dehydrogenase kinase (BCKDK), mRNA
NM_005718	Homo sapiens actin related protein 2/3 complex, subunit 4 (20 kD) (ARPC4), mRNA
NM_005717	Homo sapiens actin related protein 2/3 complex, subunit 5 (16 kD) (ARPC5), mRNA
NM_005829	Homo sapiens adaptor-related protein complex 3, sigma 2 subunit (AP3S2), mRNA
NM 005814	Homo sapiens glycoprotein A33 (transmembrane) (GPA33), mRNA

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NM_005406	Homo sapiens Rho-associated, coiled-coil containing protein kinase 1 (ROCK1), mRNA
NM_005399	Homo sapiens protein kinase, AMP-activated, beta 2 non-catalytic subunit (PRKAB2), mRNA
NM_005396	Homo sapiens pancreatic lipase-related protein 2 (PNLIPRP2), mRNA
NM 005489	Homo sapiens SH2 domain-containing 3C (SH2D3C), mRNA
NM_005479	Homo sapiens frequently rearranged in advanced T-cell lymphomas (FRAT1), mRNA
NM_005154	Homo sapiens ubiquitin specific protease 8 (USP8), mRNA
NM_005066	Homo sapiens splicing factor proline/glutamine rich (polypyrimidine tract binding protein associated) (SFPQ), mRNA
NM_005123	Homo sapiens nuclear receptor subfamily 1, group H, member 4 (NR1H4), mRNA
NM_005046	Homo sapiens kallikrein 7 (chymotryptic, stratum corneum) (KLK7), mRNA
NM_005030	Homo sapiens polo-like kinase (Drosophila) (PLK), mRNA
NM_005014	Homo sapiens osteomodulin (OMD), mRNA
NM_005003	Homo sapiens NADH dehydrogenase (ubiquinone) 1, alpha/beta subcomplex, 1 (8kD, SDAP) (NDUFAB1), mRNA
NM_004941	Homo sapiens DEAD/H (Asp-Glu-Ala-Asp/His) box polypeptide 8 (RNA helicase) (DDX8), mRNA
NM_004205	Homo sapiens ubiquitin specific protease 2 (USP2), mRNA
NM_004818	Homo sapiens prp28, U5 snRNP 100 kd protein (U5-100K), mRNA
NM 004275	Homo sapiens TRF-proximal protein (TRFP), mRNA
NM 004272	Homo sapiens Homer, neuronal immediate early gene, 1B (SYN47), mRNA
NM 004177	Homo sapiens syntaxin 3A (STX3A), mRNA
NM_004719	Homo sapiens splicing factor, arginine/serine-rich 2, interacting protein (SFRS2IP), mRNA
NM_004175	Homo sapiens small nuclear ribonucleoprotein D3 polypeptide (18kD) (SNRPD3), mRNA
NM_004592	Homo sapiens splicing factor, arginine/serine-rich 8 (suppressor-of-white-apricot homolog, Drosophila) (SFRS8), mRNA
NM_004799	Homo sapiens MAD, mothers against decapentaplegic homolog (Drosophila)
_	interacting protein, receptor activation anchor (MADHIP), transcript variant 3, mRNA
NM_004875	Homo sapiens RNA polymerase I subunit (RPA40), mRNA
NM_004292	Homo sapiens ras inhibitor (RIN1), mRNA
NM_004815	Homo sapiens PTPL1-associated RhoGAP 1 (PARG1), mRNA
NM_004772	Homo sapiens P311 protein (P311), mRNA
NM_004553	Homo sapiens NADH dehydrogenase (ubiquinone) Fe-S protein 6 (13kD) (NADH-coenzyme Q reductase) (NDUFS6), mRNA
NM_004549	Homo sapiens NADH dehydrogenase (ubiquinone) 1, subcomplex unknown, 2 (14.5kD, B14.5b) (NDUFC2), mRNA
NM_004271	Homo sapiens MD-1, RP105-associated (MD-1), mRNA
NM_004672	Homo sapiens mitogen-activated protein kinase kinase kinase 6 (MAP3K6), mRNA
NM_004828	Homo sapiens lymphocyte antigen 95 (activating NK-receptor; NK-p44) (LY95), mRNA
NM_004735	Homo sapiens leucine rich repeat (in FLII) interacting protein 1 (LRRFIP1), mRNA
NM_004811	Homo sapiens leupaxin (LPXN), mRNA
NM_004522	Homo sapiens kinesin family member 5C (KIF5C), mRNA
NM 004905	Homo sapiens anti-oxidant protein 2 (non-selenium glutathione peroxidase,
	1 orthonic protein 2 (11011-00101111111 Bramanono peroxitase,

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) 77 C 00 1770	acidic calcium-independent phospholipase A2) (KIAA0106), mRNA
NM_004770	Homo sapiens potassium voltage-gated channel, Shab-related subfamily, member 2 (KCNB2), mRNA
NM_004848	Homo sapiens basement membrane-induced gene (ICB-1), mRNA
NM_004763	Homo sapiens integrin cytoplasmic domain-associated protein 1 (ICAP-1A), transcript variant 1, mRNA
NM_004814	Homo sapiens U5 snRNP-specific 40 kDa protein (hPrp8-binding) (HPRP8BP), mRNA
NM_004839	Homo sapiens Homer, neuronal immediate early gene, 2 (HOMER-2B), mRNA
NM_004684	Homo sapiens SPARC-like 1 (mast9, hevin) (SPARCL1), mRNA
NM_004832	Homo sapiens glutathione-S-transferase like; glutathione transferase omega (GSTTLp28), mRNA
NM_004486	Homo sapiens golgi autoantigen, golgin subfamily a, 2 (GOLGA2), mRNA
NM_004125	Homo sapiens guanine nucleotide binding protein 10 (GNG10), mRNA
NM_004483	Homo sapiens glycine cleavage system protein H (aminomethyl carrier) (GCSH), mRNA
NM_004767	Homo sapiens endothelin type b receptor-like protein 2 (ET(B)R-LP-2), mRNA
NM_004440	Homo sapiens EphA7 (EPHA7), mRNA
NM_004757	Homo sapiens small inducible cytokine subfamily E, member 1 (endothelial monocyte-activating) (SCYE1), mRNA
NM_004427	Homo sapiens early development regulator 2 (polyhomeotic 2 homolog) (EDR2), mRNA
NM_004422	Homo sapiens dishevelled, dsh homolog 2 (Drosophila) (DVL2), mRNA
NM_004416	Homo sapiens deltex homolog 1 (Drosophila) (DTX1), mRNA
NM_004073	Homo sapiens cytokine-inducible kinase (CNK), mRNA
NM_004365	Homo sapiens centrin, EF-hand protein, 3 (CDC31 homolog, yeast) (CETN3), mRNA
NM_004680	Homo sapiens chromodomain protein, Y chromosome, 1 (CDY1), mRNA
NM_004291	Homo sapiens cocaine- and amphetamine-regulated transcript (CART), mRNA
NM_004330	Homo sapiens BCL2/adenovirus E1B 19kD interacting protein 2 (BNIP2), mRNA
NM_004024	Homo sapiens activating transcription factor 3 (ATF3), mRNA
NM_001177	Homo sapiens ADP-ribosylation factor-like 1 (ARL1), mRNA
NM_001545	Homo sapiens immature colon carcinoma transcript 1 (ICT1), mRNA
NM_001533	Homo sapiens heterogeneous nuclear ribonucleoprotein L (HNRPL), mRNA
NM_001509	Homo sapiens glutathione peroxidase 5 (epididymal androgen-related protein) (GPX5), transcript variant 1, mRNA
NM_001349	Homo sapiens aspartyl-tRNA synthetase (DARS), mRNA
NM_001329	Homo sapiens C-terminal binding protein 2 (CTBP2), transcript variant 1, mRNA
NM_000082	Homo sapiens Cockayne syndrome 1 (classical) (CKN1), mRNA
NM_001277	Homo sapiens choline kinase (CHK), mRNA
NM_001087	Homo sapiens angio-associated, migratory cell protein (AAMP), mRNA
NM_003999	Homo sapiens oncostatin M receptor (OSMR), mRNA
NM_003904	Homo sapiens zinc finger protein 259 (ZNF259), mRNA
NM_003385	Homo sapiens visinin-like 1 (VSNL1), mRNA
NM_003348	Homo sapiens ubiquitin-conjugating enzyme E2N (UBC13 homolog, yeast) (UBE2N), mRNA
NM_003341	Homo sapiens ubiquitin-conjugating enzyme E2E 1 (UBC4/5 homolog, yeast) (UBE2E1), mRNA
NM_003339	Homo sapiens ubiquitin-conjugating enzyme E2D 2 (UBC4/5 homolog, yeast) (UBE2D2), mRNA

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NM_003115	Homo sapiens UDP-N-acteylglucosamine pyrophosphorylase 1 (UAP1), mRNA
NM_003305	Homo sapiens transient receptor potential cation channel, subfamily C, member
	3 (TRPC3), mRNA
NM_003596	Homo sapiens tyrosylprotein sulfotransferase 1 (TPST1), mRNA
NM_003747	Homo sapiens tankyrase, TRF1-interacting ankyrin-related ADP-ribose
·	polymerase (TNKS), mRNA
NM_003569	Homo sapiens syntaxin 7 (STX7), mRNA
NM_003164	Homo sapiens syntaxin 5A (STX5A), mRNA
NM_003764	Homo sapiens syntaxin 11 (STX11), mRNA
NM_003133	Homo sapiens signal recognition particle 9kD (SRP9), mRNA
NM_003136	Homo sapiens signal recognition particle 54kD (SRP54), mRNA
NM_003131	Homo sapiens serum response factor (c-fos serum response element-binding
	transcription factor) (SRF), mRNA
NM_003795	Homo sapiens sorting nexin 3 (SNX3), mRNA
NM_003096	Homo sapiens small nuclear ribonucleoprotein polypeptide G (SNRPG), mRNA
NM_003093	Homo sapiens small nuclear ribonucleoprotein polypeptide C (SNRPC), mRNA
NM_003080	Homo sapiens sphingomyelin phosphodiesterase 2, neutral membrane (neutral
_	sphingomyelinase) (SMPD2), mRNA
NM_003059	Homo sapiens solute carrier family 22 (organic cation transporter), member 4
_	(SLC22A4), mRNA
NM_003033	Homo sapiens sialyltransferase 4A (beta-galactosidase alpha-2,3-
_	sialytransferase) (SIAT4A), mRNA
NM 003952	Homo sapiens ribosomal protein S6 kinase, 70kD, polypeptide 2 (RPS6KB2),
_	mRNA
NM 003729	Homo sapiens RTC domain containing 1 (RTCD1), mRNA
NM 002937	Homo sapiens ribonuclease, RNase A family, 4 (RNASE4), mRNA
NM 003804	Homo sapiens receptor (TNFRSF)-interacting serine-threonine kinase 1
_	(RIPK1), mRNA
NM 002898	Homo sapiens RNA binding motif, single stranded interacting protein 2
-	(RBMS2), mRNA
NM_002886	Homo sapiens RAP2B, member of RAS oncogene family (RAP2B), mRNA
NM_003953	Homo sapiens myelin protein zero-like 1 (MPZL1), mRNA
NM_002809	Homo sapiens proteasome (prosome, macropain) 26S subunit, non-ATPase, 3
-	(PSMD3), mRNA
NM_002771	Homo sapiens protease, serine, 3 (trypsin 3) (PRSS3), mRNA
NM_002757	Homo sapiens mitogen-activated protein kinase kinase 5 (MAP2K5), mRNA
NM_002754	Homo sapiens mitogen-activated protein kinase 13 (MAPK13), mRNA
NM_003668	Homo sapiens mitogen-activated protein kinase-activated protein kinase 5
-	(MAPKAPK5), mRNA
NM_002718	Homo sapiens protein phosphatase 2 (formerly 2A), regulatory subunit B" (PR
	72), alpha isoform and (PR 130), beta isoform (PPP2R3), mRNA
NM_003622	Homo sapiens PTPRF interacting protein, binding protein 1 (liprin beta 1)
	(PPFIBP1), mRNA
NM_003626	Homo sapiens protein tyrosine phosphatase, receptor type, f polypeptide
	(PTPRF), interacting protein (liprin), alpha 1 (PPFIA1), mRNA
NM_002689	Homo sapiens polymerase (DNA-directed), alpha (70kD) (POLA2), mRNA
NM 002685	Homo sapiens polymyositis/scleroderma autoantigen 2 (100kD) (PMSCL2),
-	mRNA
NM 003876	Homo sapiens putative receptor protein (PMI), mRNA
NM 002670	Homo sapiens plastin 1 (I isoform) (PLS1), mRNA
NM 002664	Homo sapiens pleckstrin (PLEK), mRNA
NM 003559	Homo sapiens phosphatidylinositol-4-phosphate 5-kinase, type II, beta
	The brookstand mores brookstand a strength of the religious

<u> </u>	(PIP5K2B), mRNA
NM_003629	Homo sapiens phosphoinositide-3-kinase, regulatory subunit, polypeptide 3 (p55,
	gamma) (PIK3R3), mRNA
NM_002649	Homo sapiens phosphoinositide-3-kinase, catalytic, gamma polypeptide
	(PIK3CG), mRNA
NM_002624	Homo sapiens prefoldin 5 (PFDN5), mRNA
NM_003846	Homo sapiens peroxisomal biogenesis factor 11B (PEX11B), mRNA
NM_002617	Homo sapiens peroxisome biogenesis factor 10 (PEX10), mRNA
NM_002611	Homo sapiens pyruvate dehydrogenase kinase, isoenzyme 2 (PDK2), mRNA
NM_000923	Homo sapiens phosphodiesterase 4C; cAMP-specific (phosphodiesterase E1 dunce homolog, Drosophila) (PDE4C), mRNA
NM 002599	Homo sapiens phosphodiesterase 2A, cGMP-stimulated (PDE2A), mRNA
NM 002504	Homo sapiens nuclear transcription factor, X-box binding 1 (NFX1), mRNA
NM_002482	Homo sapiens nuclear autoantigenic sperm protein (histone-binding) (NASP), mRNA
NM_003826	Homo sapiens N-ethylmaleimide-sensitive factor attachment protein, gamma (NAPG), mRNA
NM 002465	Homo sapiens myosin binding protein C, slow type (MYBPC1), mRNA
NM 002461	Homo sapiens mevalonate (diphospho) decarboxylase (MVD), mRNA
NM_003676	Homo sapiens degenerative spermatocyte homolog, lipid desaturase (Drosophila)
	(DEGS), mRNA
NM_002307	Homo sapiens lectin, galactoside-binding, soluble, 7 (galectin 7) (LGALS7),
	mRNA
NM_002271	Homo sapiens karyopherin (importin) beta 3 (KPNB3), mRNA
NM_002270	Homo sapiens karyopherin (importin) beta 2 (KPNB2), mRNA
NM_002214	Homo sapiens integrin, beta 8 (ITGB8), mRNA
NM_002204	Homo sapiens integrin, alpha 3 (antigen CD49C, alpha 3 subunit of VLA-3 receptor) (ITGA3), transcript variant a, mRNA
NM 001560	Homo sapiens interleukin 13 receptor, alpha 1 (IL13RA1), mRNA
NM_002163	Homo sapiens interferon consensus sequence binding protein 1 (ICSBP1), mRNA
NR4 000156	Homo sapiens heat shock 60kD protein 1 (chaperonin) (HSPD1), mRNA
NM_002156	Homo sapiens hippocalcin-like 1 (HPCAL1), mRNA
NM_002149 NM_003947	Homo sapiens huntingtin-associated protein interacting protein (duo) (HAPIP), mRNA
NM_003665	Homo sapiens ficolin (collagen/fibrinogen domain containing) 3 (Hakata antigen) (FCN3), mRNA
NM_000842	Homo sapiens glutamate receptor, metabotropic 5 (GRM5), mRNA
NM 002053	Homo sapiens guanylate binding protein 1, interferon-inducible, 67kD (GBP1),
14147_005022	mRNA
NM_001482	Homo sapiens glycine amidinotransferase (L-arginine:glycine
	amidinotransferase) (GATM), mRNA
NM 002044	Homo sapiens galactokinase 2 (GALK2), mRNA
NM_001417	Homo sapiens eukaryotic translation initiation factor 4B (EIF4B), mRNA
NM_003758	Homo sapiens eukaryotic translation initiation factor 3, subunit 1 (alpha, 35kD) (EIF3S1), mRNA
NM_001404	Homo sapiens eukaryotic translation elongation factor 1 gamma (EEF1G), mRNA
NM_001960	Homo sapiens eukaryotic translation elongation factor 1 delta (guanine nucleotide exchange protein) (EEF1D), mRNA
NM 003792	Homo sapiens endothelial differentiation-related factor 1 (EDF1), mRNA
NM_003974	Homo sapiens docking protein 2, 56kD (DOK2), mRNA

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NM_003586	Homo sapiens double C2-like domains, alpha (DOC2A), mRNA
NM_001883	Homo sapiens corticotropin releasing hormone receptor 2 (CRHR2), mRNA
NM_001873	Homo sapiens carboxypeptidase E (CPE), mRNA
NM_001782	Homo sapiens CD72 antigen (CD72), mRNA
NM_001762	Homo sapiens chaperonin containing TCP1, subunit 6A (zeta 1) (CCT6A), mRNA
NM_003716	Homo sapiens Ca2+-dependent activator protein for secretion (CADPS), mRNA
NM_003986	Homo sapiens butyrobetaine (gamma), 2-oxoglutarate dioxygenase (gamma-butyrobetaine hydroxylase) 1 (BBOX1), mRNA
NM 001674	Homo sapiens activating transcription factor 3 (ATF3), mRNA
NM 001173	Homo sapiens Rho GTPase activating protein 5 (ARHGAP5), mRNA
NM 025065	Homo sapiens RNA processing factor 1 (RPF1), mRNA
NM 024907	Homo sapiens F-box protein FBG4 (FBG4), mRNA
NM 025194	Homo sapiens inositol 1,4,5-trisphosphate 3-kinase C (ITPKC), mRNA
NM_014203	Homo sapiens adaptor-related protein complex 2, alpha 1 subunit (AP2A1),
NM_130786	mRNA
NM 031482	Homo sapiens alpha-1-B glycoprotein (A1BG), mRNA Homo sapiens hypothesical protein DVE7-58-G0418 (DVE7D58-G0418) DVA
NM 015419	Homo sapiens hypothetical protein DKFZp586I0418 (DKFZP586I0418), mRNA
NM 015683	Homo sapiens adlican (DKFZp564I1922), mRNA
NM 015638	Homo sapiens hypothetical protein (CLONE24945), mRNA
NM_080737	Homo sapiens chromosome 20 open reading frame 188 (C20orf188), mRNA
NM 080723	Homo sapiens synaptotagmin-like 4 (granuphilin-a) (SYTLA), mRNA
NM 080678	Homo sapiens vesicular membrane protein p24 (VMP), mRNA
NM_080668	Homo sapiens NEDD8-conjugating enzyme (NCE2), mRNA
	Homo sapiens similar to RIKEN cDNA 2610036L13 (MGC16386), mRNA
NM_080666	Homo sapiens similar to RIKEN cDNA 2600001A11 gene (LOC112840), mRNA
NM_080663	Homo sapiens similar to RIKEN cDNA 4933424N09 gene (MGC16943), mRNA
NM_080661	Homo sapiens similar to RIKEN cDNA 0610008P16 gene (MGC15937), mRNA
NM_080658	Homo sapiens similar to RIKEN cDNA 0610006H10 gene (MGC9740), mRNA
NM_080656	Homo sapiens similar to RIKEN cDNA A430101B06 gene (MGC13017), mRNA
NM_080651	Homo sapiens similar to RIKEN cDNA 1810038N03 gene (MGC9890), mRNA
NM_080650	Homo sapiens similar to RIKEN cDNA 5730421E18 gene (MGC14798), mRNA
NM_080604	Homo sapiens tight junction protein 4 (peripheral) (TJP4), mRNA
NM_080552	Homo sapiens vesicular inhibitory amino acid transporter (VIAAT), mRNA
NM_080429	Homo sapiens aquaporin 10 (AQP10), mRNA
NM_018897	Homo sapiens axonemal dynein heavy chain 7 (DNAH7), mRNA
NM_015570	Homo sapiens autism-related protein 1 (KIAA0442), mRNA
NM_015132	Homo sapiens sorting nexin 13 (SNX13), mRNA
NM_022457	Homo sapiens similar to constitutive photomorphogenic protein 1 (Arabidopsis) (FLJ10416), mRNA
NM_030658	Homo sapiens putative ankyrin-repeat containing protein (DKFZP564D166), mRNA
NM_058229	Homo sapiens F-box only protein 32 (FBXO32), mRNA
NM_058188	Homo sapiens chromosome 21 open reading frame 67 (C21orf67), mRNA
NM_058187	Homo sapiens chromosome 21 open reading frame 63 (C21orf63), mRNA
NM_058171	Homo sapiens ING1-like tumor suppressor protein (ING1-like), mRNA
NM_058167	Homo sapiens ubiquitin conjugating enzyme 6 (Ubc6p), mRNA
NM 015242	Homo sapiens centaurin, delta 2 (CENTD2), mRNA
NM_054114	Homo sapiens hypothetical protein FLJ32631 (FLJ32631), mRNA
NM 054111	TI protecti i Labout (1 Labout), mutat
INIMI ODALITI I	Homo sapiens inositol hexaphosphate kinase 3 (IHPK3), mRNA

NM_054108	Homo sapiens H-rev107-like protein 5 (HRLP5), mRNA
NM_020794	Homo sapiens densin-180 (KIAA1365), mRNA
NM_054032	Homo sapiens G protein-coupled receptor MRGX4 (MRGX4), mRNA
NM_054031	Homo sapiens G protein-coupled receptor MRGX3 (MRGX3), mRNA
NM_054030	Homo sapiens G protein-coupled receptor MRGX2 (MRGX2), mRNA
NM_054023	Homo sapiens uteroglobin-related protein 1 (UGRP1), mRNA
NM_054024	Homo sapiens melanoma inhibitory activity protein 2 (MIA2), mRNA
NM_031946	Homo sapiens centaurin, gamma 3 (CENTG3), mRNA
NM_052860	Homo sapiens kruppel-like zinc finger protein (ZNF300), mRNA
NM_053054	Homo sapiens cation channel of sperm (CATSPER), mRNA
NM_053053	Homo sapiens SPT3-associated factor 42 (STAF42), mRNA
NM_053048	Homo sapiens hypothetical protein MGC16384 (MGC16384), mRNA
NM_053047	Homo sapiens hypothetical protein MGC16063 (MGC16063), mRNA
NM_053040	Homo sapiens PNAS-123 (LOC85028), mRNA
NM_053039	Homo sapiens UDP glycosyltransferase 2 family, polypeptide B28 (UGT2B28), mRNA
NM_053001	Homo sapiens odd-skipped-related 2A protein (OSR2), mRNA
NM_052997	Homo sapiens breast cancer antigen NY-BR-1 (NY-BR-1), mRNA
NM_052971	Homo sapiens liver-expressed antimicrobial peptide 2 (LEAP-2), mRNA
NM_052956	Homo sapiens medium-chain acyl-CoA synthetase (MACS1), mRNA
NM_052942	Homo sapiens guanylate binding protein 5 (GBP5), mRNA
NM_052931	Homo sapiens activating NK receptor (KALI), mRNA
NM_052879	Homo sapiens c-Mpl binding protein (LOC113251), mRNA
NM_030928	Homo sapiens DNA replication factor (CDT1), mRNA
NM_025185	Homo sapiens putative ankyrin-repeat containing protein (DKFZP564D166), mRNA
NM_015179	Homo sapiens KIAA0690 protein (KIAA0690), mRNA
NM_033626	Homo sapiens JM11 protein (JM11), mRNA
NM_022735	Homo sapiens golgi phosphoprotein 1 (GOLPH1), mRNA
NM_033547	Homo sapiens hypothetical gene MGC16733 similar to CG12113 (MGC16733), mRNA
NM_032268	Homo sapiens nerve injury gene 283 (NIN283), mRNA
NM_016167	Homo sapiens retinoic acid repressible protein (RARG-1), mRNA
NM_033414	Homo sapiens hypothetical protein MGC17552 (MGC17552), mRNA
NM_016336	Homo sapiens non-canonical ubquitin conjugating enzyme 1 (NCUBE1), mRNA
NM_033317	Homo sapiens hypothetical gene ZD52F10 (ZD52F10), mRNA
NM_033266	Homo sapiens ER to nucleus signalling 2 (ERN2), mRNA
NM_031955	Homo sapiens NYD-SP12 protein (NYD-SP12), mRNA
NM_033210	Homo sapiens hypothetical protein FLJ14855 (FLJ14855), mRNA
NM_033211	Homo sapiens hypothetical gene supported by AF038182; BC009203 (LOC90355), mRNA
NM_033194	Homo sapiens small heat shock protein B9 (HspB9), mRNA
NM_032122	Homo sapiens dystrobrevin binding protein 1 (DTNBP1), mRNA
NM_020405	Homo sapiens tumor endothelial marker 7 precursor (TEM7), mRNA
NM_033115	Homo sapiens hypothetical protein MGC16169 (MGC16169), mRNA
NM_033117	Homo sapiens hypothetical protein MGC2734 (MGC2734), mRNA
NM_033103	Homo sapiens rhophilin-like protein (LOC85415), mRNA
NM_033035	Homo sapiens thymic stromal lymphopoietin (TSLP), mRNA
NM_014001	Homo sapiens golgi associated, gamma adaptin ear containing, ARF binding protein 3 (GGA3), mRNA
NM 015149	Homo sapiens RalGDS-like gene (RGL), mRNA
NM_032937	Homo sapiens AD038 (LOC85026), mRNA

NM_032932	Homo sapiens hypothetical protein MGC11316 (MGC11316), mRNA
NM_032930	Homo sapiens hypothetical protein MGC13040 (MGC13040), mRNA
NM_032918	Homo sapiens RAS-like, estrogen-regulated, growth-inhibitor (RERG), mRNA
NM_032916	Homo sapiens hypothetical protein MGC16279 (MGC16279), mRNA
NM_032907	Homo sapiens hypothetical protein MGC14421 (MGC14421), mRNA
NM_032904	Homo sapiens hypothetical protein MGC14433 (MGC14433), mRNA
NM_032900	Homo sapiens hypothetical protein MGC14258 (MGC14258), mRNA
NM_032895	Homo sapiens hypothetical protein MGC14376 (MGC14376), mRNA
NM_032888	Homo sapiens KIAA1870 protein (KIAA1870), mRNA
NM_032886	Homo sapiens hypothetical protein MGC15912 (MGC15912), mRNA
NM_032884	Homo sapiens hypothetical protein MGC15882 (MGC15882), mRNA
NM_032876	Homo sapiens hypothetical protein MGC15563 (MGC15563), mRNA
NM_032875	Homo sapiens hypothetical protein MGC15482 (MGC15482), mRNA
NM_032874	Homo sapiens hypothetical protein MGC15438 (MGC15438), mRNA
NM_032872	Homo sapiens NADPH oxidase-related, C2 domain-containing protein (JFC1), mRNA
NM_032871	Homo sapiens tumor necrosis factor receptor superfamily, member 19-like (TNFRSF19L), mRNA
NM_032866	Homo sapiens hypothetical protein FLJ14957 (FLJ14957), mRNA
NM_032860	Homo sapiens hypothetical protein FLJ14909 (FLJ14909), mRNA
NM_032858	Homo sapiens hypothetical protein FLJ14904 (FLJ14904), mRNA
NM_032852	Homo sapiens AUT-like 1, cysteine endopeptidase (S. cerevisiae) (AUTL1), mRNA
NM 032848	Homo sapiens hypothetical protein FLJ14827 (FLJ14827), mRNA
NM_032845	Homo sapiens hypothetical protein FLJ14816 (FLJ14816), mRNA
NM 032835	Homo sapiens hypothetical protein FLJ14761 (FLJ14761), mRNA
NM 032824	Homo sapiens hypothetical protein FLJ14681 (FLJ14681), mRNA
NM 032823	Homo sapiens hypothetical protein FLJ14675 (FLJ14675), mRNA
NM 032822	Homo sapiens hypothetical protein FLJ14668 (FLJ14668), mRNA
NM 032818	Homo sapiens hypothetical protein FLJ14642 (FLJ14642), mRNA
NM 032804	Homo sapiens hypothetical protein FLJ14547 (FLJ14547), mRNA
NM_032795	Homo sapiens hypothetical protein FLJ14494 (FLJ14494), mRNA
NM_032783	Homo sapiens hypothetical protein FLJ14431 (FLJ14431), mRNA
NM_032766	Homo sapiens hypothetical protein MGC16179 (MGC16179), mRNA
NM 032763	Homo sapiens hypothetical protein MGC16142 (MGC16142), mRNA
NM_032756	Homo sapiens hypothetical protein MGC15668 (MGC15668), mRNA
NM_032744	Homo sapiens hypothetical protein MGC12335 (MGC12335), mRNA
NM_032738	Homo sapiens hypothetical protein MGC4595 (MGC4595), mRNA
NM_032723	Homo sapiens hypothetical protein MGC12760 (MGC12760), mRNA
NM_032720	Homo sapiens hypothetical protein MGC10724 (MGC10724), mRNA
NM_032715	Homo sapiens hypothetical protein MGC4643 (MGC4643), mRNA
NM_032712	Homo sapiens hypothetical protein MGC13170 (MGC13170), mRNA
NM 032711	Homo sapiens hypothetical protein MGC13090 (MGC13090), mRNA
NM 032706	Homo sapiens hypothetical protein MGC12966 (MGC12966), mRNA
NM_032705	Homo sapiens hypothetical protein MGC14801 (MGC14801), mRNA
NM 032694	Homo sapiens hypothetical protein MGC12935 (MGC12935), mRNA
NM_032693	Homo sapiens hypothetical protein MGC10646 (MGC10646), mRNA
NM_032681	Homo sapiens hypothetical protein MGC10977 (MGC10977), mRNA
NM_032678	Homo sapiens hypothetical protein MGC3413 (MGC3413), mRNA
NM_032667	Homo sapiens hypothetical protein MGC4694 (MGC4694), mRNA
NM_032661	Homo sapiens hypothetical protein MGC5139 (MGC5139), mRNA
NM 032634	Homo sapiens hypothetical protein MGC3079 (MGC3079), mRNA
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NM 032601 NM 032596 NM 032593 NM 032586 NM 032582 NM 032580 NM 032574 NM 032558	Homo sapiens hypothetical protein MGC2641 (MGC2641), mRNA Homo sapiens methylmalonyl CoA epimerase (MCEE), mRNA Homo sapiens testes development-related NYD-SP22 (NYD-SP22), mRNA Homo sapiens PKCI-1-related HIT protein (HIT-17), mRNA Homo sapiens testis transcript Y 8 (TTY8), mRNA Homo sapiens ubiquitin specific protease (NY-REN-60), mRNA Homo sapiens hairy and enhancer of split 7 (Drosophila) (HES7), mRNA Homo sapiens dpy-30-like protein (LOC84661), mRNA Homo sapiens hypothetical protein FLJ14753 (FLJ14753), mRNA Homo sapiens HP43.8KD protein (HP43.8KD), mRNA Homo sapiens putative purinergic receptor (FKSG79), mRNA Homo sapiens cryptic gene (CRYPTIC), mRNA Homo sapiens Mov10, Moloney leukemia virus 10, homolog (mouse) (MOV10),
NM 032596 NM 032593 NM 032586 NM 032582 NM 032580 NM 032574 NM 032558	Homo sapiens testes development-related NYD-SP22 (NYD-SP22), mRNA Homo sapiens PKCI-1-related HIT protein (HIT-17), mRNA Homo sapiens testis transcript Y 8 (TTY8), mRNA Homo sapiens ubiquitin specific protease (NY-REN-60), mRNA Homo sapiens hairy and enhancer of split 7 (Drosophila) (HES7), mRNA Homo sapiens dpy-30-like protein (LOC84661), mRNA Homo sapiens hypothetical protein FLJ14753 (FLJ14753), mRNA Homo sapiens HP43.8KD protein (HP43.8KD), mRNA Homo sapiens putative purinergic receptor (FKSG79), mRNA Homo sapiens cryptic gene (CRYPTIC), mRNA Homo sapiens Mov10, Moloney leukemia virus 10, homolog (mouse) (MOV10),
NM_032593 NM_032586 NM_032582 NM_032580 NM_032574 NM_032558	Homo sapiens PKCI-1-related HTT protein (HIT-17), mRNA Homo sapiens testis transcript Y 8 (TTY8), mRNA Homo sapiens ubiquitin specific protease (NY-REN-60), mRNA Homo sapiens hairy and enhancer of split 7 (Drosophila) (HES7), mRNA Homo sapiens dpy-30-like protein (LOC84661), mRNA Homo sapiens hypothetical protein FLJ14753 (FLJ14753), mRNA Homo sapiens HP43.8KD protein (HP43.8KD), mRNA Homo sapiens putative purinergic receptor (FKSG79), mRNA Homo sapiens cryptic gene (CRYPTIC), mRNA Homo sapiens Mov10, Moloney leukemia virus 10, homolog (mouse) (MOV10),
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NM 032582 NM 032580 NM 032574 NM 032558	Homo sapiens ubiquitin specific protease (NY-REN-60), mRNA Homo sapiens hairy and enhancer of split 7 (Drosophila) (HES7), mRNA Homo sapiens dpy-30-like protein (LOC84661), mRNA Homo sapiens hypothetical protein FLJ14753 (FLJ14753), mRNA Homo sapiens HP43.8KD protein (HP43.8KD), mRNA Homo sapiens putative purinergic receptor (FKSG79), mRNA Homo sapiens cryptic gene (CRYPTIC), mRNA Homo sapiens Mov10, Moloney leukemia virus 10, homolog (mouse) (MOV10),
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	Homo sapiens cryptic gene (CRYPTIC), mRNA Homo sapiens Mov10, Moloney leukemia virus 10, homolog (mouse) (MOV10),
NM_032553	Homo sapiens cryptic gene (CRYPTIC), mRNA Homo sapiens Mov10, Moloney leukemia virus 10, homolog (mouse) (MOV10),
	mRNA
	Homo sapiens hypothetical protein MGC2629 (MGC2629), mRNA
	Homo sapiens cerebral protein-4 (HUCEP-4), mRNA
	Homo sapiens hypothetical protein HH114 (HH114), mRNA
	Homo sapiens zinc finger protein (LOC84524), mRNA
	Homo sapiens hypothetical protein GL009 (GL009), mRNA
	Homo sapiens actin related protein M1 (ARPM1), mRNA
	Homo sapiens dynactin 4 (MGC3248), mRNA
	Homo sapiens MEGF11 protein (MEGF11), mRNA
	Homo sapiens hypothetical protein FLJ21673 (FLJ21673), mRNA
	Homo sapiens putative nuclear protein ORF1-FL49 (ORF1-FL49), mRNA
	Homo sapiens esophageal cancer related gene 4 protein (ECRG4), mRNA
	Homo sapiens cylindromatosis (turban tumor syndrome) (CYLD), mRNA
	Homo sapiens hypothetical protein MGC12536 (MGC12536), mRNA
	Homo sapiens hypothetical protein FLJ23183 (FLJ23183), mRNA
	Homo sapiens hypothetical protein MGC16186 (MGC16186), mRNA
	Homo sapiens hypothetical protein MGC15435 (MGC15435), mRNA
	Homo sapiens hypothetical protein MGC10744 (MGC10744), mRNA
	Homo sapiens hypothetical protein MGC13250 (MGC13250), mRNA
	Homo sapiens hypothetical protein MGC13045 (MGC13045), mRNA
	Homo sapiens hypothetical protein MGC12992 (MGC12992), mRNA
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	Homo sapiens hypothetical protein MGC14817 (MGC14817), mRNA
	Homo sapiens hypothetical protein MGC4248 (MGC4248), mRNA
	Homo sapiens hypothetical protein MGC2993 (MGC2993), mRNA
	Homo sapiens hypothetical protein MGC11102 (MGC11102), mRNA
NM_032324	Homo sapiens hypothetical protein MGC13186 (MGC13186), mRNA
	Homo sapiens hypothetical protein MGC13102 (MGC13102), mRNA
	Homo sapiens hypothetical protein MGC13007 (MGC13007), mRNA
	Homo sapiens hypothetical protein MGC12945 (MGC12945), mRNA
	Homo sapiens hypothetical protein MGC12943 (MGC12943), mRNA
	Homo sapiens hypothetical protein MGC12936 (MGC12936), mRNA
NM_032305	Homo sapiens hypothetical protein MGC3200 (MGC3200), mRNA
	Homo sapiens hypothetical protein DKFZp761J1523 (DKFZp761J1523), mRNA
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NM_032290 1	Homo sapiens hypothetical protein DKFZp761C121 (DKFZp761C121), mRNA
	Homo sapiens hypothetical protein DKFZp761B1514 (DKFZp761B1514), mRNA
	Homo sapiens hypothetical protein DKFZp586C1924 (DKFZp586C1924),
	Examo suprens hypothetical protein DATZp360C1324 (DATZp380C1924),

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NM_032299	Homo sapiens hypothetical protein MGC2714 (MGC2714), mRNA
	Homo sapiens hypothetical protein DKFZp434E169 (DKFZp434E169), mRNA
	Homo sapiens hypothetical protein DKFZp434D177 (DKFZp434D177), mRNA
-	Homo sapiens hypothetical protein DKFZp434N0650 (DKFZp434N0650), mRNA
	Homo sapiens hypothetical protein DKFZp434P2235 (DKFZp434P2235), mRNA
	Homo sapiens hypothetical protein DKFZp434G0920 (DKFZp434G0920), mRNA
NM 032250	Homo sapiens hypothetical protein DKFZp434A171 (DKFZp434A171), mRNA
NM_032249	Homo sapiens hypothetical protein DKFZp434F1819 (DKFZp434F1819), mRNA
	Homo sapiens hypothetical protein DKFZp434F1719 (DKFZp434F1719), mRNA
	Homo sapiens hypothetical protein DKFZp434J0617 (DKFZp434J0617), mRNA
	Homo sapiens hypothetical protein DKFZp434I1916 (DKFZp434I1916), mRNA
	Homo sapiens hypothetical protein FLJ22427 (FLJ22427), mRNA
NM_032209	Homo sapiens hypothetical protein FLJ21777 (FLJ21777), mRNA
NM_032193	Homo sapiens hypothetical protein FLJ20974 (FLJ20974), mRNA
	Homo sapiens hypothetical protein FLJ13193 (FLJ13193), mRNA
NM_032167	Homo sapiens hypothetical protein FLJ12363 (FLJ12363), mRNA
NM_032161	Homo sapiens KIAA1870 protein (KIAA1870), mRNA
NM_032154	Homo sapiens MBLR protein (MBLR), mRNA
NM_032151	Homo sapiens hypothetical protein DKFZp566K1946 (DKFZP566K1946), mRNA
NM_032148	Homo sapiens hypothetical protein DKFZp434K0427 (DKFZP434K0427), mRNA
	Homo sapiens hypothetical protein DKFZp434L0718 (DKFZP434L0718), mRNA
	Homo sapiens hypothetical protein DKFZp434E2318 (DKFZP434E2318), mRNA
	Homo sapiens hypothetical protein DKFZp434L1717 (DKFZP434L1717), mRNA
	Homo sapiens hypothetical protein DKFZp564D0478 (DKFZP564D0478), mRNA
	Homo sapiens hypothetical protein DKFZp564O0523 (DKFZP564O0523), mRNA
	Homo sapiens ninein (GSK3B interacting protein) (NIN), mRNA
	Homo sapiens hypothetical protein DKFZp762I166 (DKFZP762I166), mRNA
	Homo sapiens hypothetical protein DKFZp762L0311 (DKFZp762L0311), mRNA
NM_015630	Homo sapiens DKFZP566F2124 protein (DKFZP566F2124), mRNA
NM_015621	Homo sapiens DKFZP434C171 protein (DKFZP434C171), mRNA
	Homo sapiens DKFZP434D146 protein (DKFZP434D146), mRNA
NM_015496	Homo sapiens DKFZP434I116 protein (DKFZP434I116), mRNA
NM_015471	Homo sapiens DKFZP566O1646 protein (DC8), mRNA
NM_015453	Homo sapiens DKFZP434F091 protein (DKFZP434F091), mRNA
NM_015023	Homo sapiens KIAA1037 protein (KIAA1037), mRNA
	Homo sapiens KIAA1049 protein (KIAA1049), mRNA
	220 Mo Dapiens IEE H11049 protein (IEE H110 19); IMC411
NM_032042	Homo sapiens hypothetical protein DKFZp564D172 (DKFZP564D172), mRNA

NM_032030	Homo sapiens FKSG83 (FKSG83), mRNA
NM_032028	Homo sapiens serine/threonine kinase FKSG81 (FKSG81), mRNA
NM_032025	Homo sapiens CDA02 protein (CDA02), mRNA
NM_032021	Homo sapiens AD031 protein (AD031), mRNA
NM_031944	Homo sapiens Mix-like homeobox protein 1 (MILD1), mRNA
NM_031920	Homo sapiens ARG99 protein (ARG99), mRNA
NM_031480	Homo sapiens hypothetical protein AD034 (AD034), mRNA
NM_031478	Homo sapiens hypothetical protein DKFZp434I2117 (DKFZP434I2117), mRNA
NM_031477	Homo sapiens hypothetical protein MGC10500 (MGC10500), mRNA
NM_031476	Homo sapiens hypothetical protein DKFZp434B044 (DKFZP434B044), mRNA
NM_031472	Homo sapiens hypothetical protein MGC11134 (MGC11134), mRNA
NM 031471	Homo sapiens hypothetical protein MGC10966 (MGC10966), mRNA
NM_031457	Homo sapiens membrane-spanning 4-domains, subfamily A, member 8B
	(MS4A8B), mRNA
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NM_031443	Homo sapiens hypothetical protein MGC4607 (MGC4607), mRNA
NM_031438	Homo sapiens hypothetical protein DKFZp761I172 (DKFZP761I172), mRNA
NM_031434	Homo sapiens hypothetical protein MGC5442 (MGC5442), mRNA
NM_031418	Homo sapiens chromosome 11 open reading frame 25 (C11orf25), mRNA
NM_015497	Homo sapiens DKFZP564G2022 protein (DKFZP564G2022), mRNA
NM_031306	Homo sapiens hypothetical protein DKFZp564B1023 (DKFZP564B1023), mRNA
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NM_031291	Homo sapiens hypothetical protein DKFZp434N1235 (DKFZP434N1235), mRNA
NM_031290	Homo sapiens hypothetical protein DKFZp434K1172 (DKFZP434K1172), mRNA
NM_031270	Homo sapiens PRO1596 protein (PRO1596), mRNA
NM 031268	Homo sapiens PRO0461 protein (PRO0461), mRNA
NM_031217	Homo sapiens hypothetical protein DKFZp434G2226 (DKFZP434G2226), mRNA
NM 013358	Homo sapiens peptidylarginine deiminase type I (hPAD-colony10), mRNA
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NM 030954	Homo sapiens hypothetical protein DKFZp564A022 (DKFZP564A022), mRNA
NM_030953	Homo sapiens hypothetical protein DKFZp761E2110 (DKFZP761E2110), mRNA
NM 030941	Homo sapiens exonuclease NEF-sp (LOC81691), mRNA
NM_030939	Homo sapiens hypothetical protein FLJ12619 (FLJ12619), mRNA
NM_030938	Homo sapiens likely ortholog of rat vacuole membrane protein 1 (VMP1), mRNA
NM 030932	Homo sapiens diaphanous homolog 3 (Drosophila) (DIAPH3), mRNA
NM 030927	Homo sapiens hypothetical protein MGC11352 (MGC11352), mRNA
NM 030925	Homo sapiens hypothetical protein FLJ12577 (FLJ12577), mRNA
NM 030918	Homo sapiens hypothetical protein My014 (MY014), mRNA
NM_030911	Homo sapiens protein kinase NYD-SP15 (NYD-SP15), mRNA
NM 030899	Homo sapiens hypothetical protein FLJ23407 (FLJ23407), mRNA
NM 018657	Homo sapiens myoneurin (MYNN), mKNA
NM_018657 NM_030818	Homo sapiens myoneurin (MYNN), mRNA Homo sapiens hypothetical protein MGC10471 (MGC10471), mRNA
NM_030818	Homo sapiens hypothetical protein MGC10471 (MGC10471), mRNA
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	mRNA
NM 030802	Homo sapiens C/EBP-induced protein (LOC81558), mRNA
NM 030800	Homo sapiens hypothetical protein DKFZp564O1664 (DKFZP564O1664),
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NM 030799	Homo sapiens hypothetical protein AF140225 (AF140225), mRNA
NM 030793	Homo sapiens hypothetical protein SP329 (SP329), mRNA
NM 030792	Homo sapiens hypothetical protein PP1665 (PP1665), mRNA
	Homo sapiens folate transporter/carrier (LOC81034), mRNA
NM 030780	Homo sapiens solute carrier family 38, member 1 (SLC38A1), mRNA
NM_030674	Homo sapiens hypothetical protein FLJ10312 (FLJ10312), mRNA
NM_030672	Homo sapiens hypothetical protein FLJ12729 (FLJ12729), mRNA
NM_024947 NM_024963	Homo sapiens hypothetical protein FLJ11467 (FLJ11467), mRNA
	Homo sapiens hypothetical protein DKFZp434M0331 (DKFZp434M0331),
NM_017600	mRNA
NM 030652	Homo sapiens NG3 protein (NG3), mRNA
NM 030651	Homo sapiens chromosome 6 open reading frame 31 (C6orf31), mRNA
NM 020444	Homo sapiens KIAA1191 protein (KIAA1191), mRNA
NM_020444 NM_024055	Homo sapiens KIAAT191 protein (KIAAT191), inklyA Homo sapiens hypothetical protein MGC5499 (MGC5499), mRNA
NM 025154	Homo sapiens KIAA0810 protein (KIAA0810), mRNA
	Homo sapiens novel protein (HSNOV1), mRNA
NM_017515 NM_024924	Homo sapiens hypothetical protein (HSNOVI), mkNA Homo sapiens hypothetical protein FLJ12985 (FLJ12985), mRNA
NM_030579	Homo sapiens cytochrome b5 outer mitochondrial membrane precursor (CYB5-
14141_030379	M), mRNA
NM 022068	Homo sapiens hypothetical protein FLJ23403 (FLJ23403), mRNA
NM_025179	Homo sapiens plexin A2 (PLXNA2), mRNA
NM 014033	Homo sapiens DKFZP586A0522 protein (DKFZP586A0522), mRNA
NM_006468	Homo sapiens polymerase (RNA) III (DNA directed) (62kD) (RPC62), mRNA
NM 025263	Homo sapiens CAT56 protein (CAT56), mRNA
NM 025262	Homo sapiens G5C protein (G5C), mRNA
NM 025261	Homo sapiens G6C protein (G6C), mRNA
NM_025260	Homo sapiens G6B protein (G6B), mRNA
NM_025259	Homo sapiens NG23 protein (NG23), mRNA
NM 025258	Homo sapiens NG37 protein (G7C), mRNA
NM_025231	Homo sapiens hypothetical protein FLJ22191 (FLJ22191), mRNA
NM 025226	Homo sapiens MSTP032 protein (MSTP032), mRNA
NM 025211	Homo sapiens protein kinase anchoring protein GKAP42 (GKAP42), mRNA
NM 025201	Homo sapiens hypothetical protein PP1628 (PP1628), mRNA
NM_025192	Homo sapiens hypothetical protein FLJ23071 (FLJ23071), mRNA
NM_025188	Homo sapiens hypothetical protein FLJ13181 (FLJ13181), mRNA
NM 025174	Homo sapiens hypothetical protein FLJ23040 (FLJ23040), mRNA
NM_025165	Homo sapiens hypothetical protein FLJ22637 (FLJ22637), mRNA
NM_025160	Homo sapiens hypothetical protein FLJ21016 (FLJ21016), mRNA
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NM_025151	Homo sapiens hypothetical protein FLJ22622 (FLJ22622), mRNA
NM 025149	Homo sapiens hypothetical protein FLJ20920 (FLJ20920), mRNA
NM 025144	Homo sapiens hypothetical protein FLJ22670 (FLJ22670), mRNA
NM 025138	Homo sapiens hypothetical protein FLJ12661 (FLJ12661), mRNA
NM 025126	Homo sapiens ring finger protein 34 (RNF34), mRNA
NM_025125	Homo sapiens hypothetical protein FLJ13263 (FLJ13263), mRNA
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NM_025077	Homo sapiens hypothetical protein FLJ13949 (FLJ13949), mRNA
NM_025076	Homo sapiens hypothetical protein FLJ23591 (FLJ23591), mRNA
NM_025072	Homo sapiens chromosome 9 open reading frame 15 (C9orf15), mRNA
NM_025070	Homo sapiens hypothetical protein FLJ22242 (FLJ22242), mRNA
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NM_025055	Homo sapiens hypothetical protein FLJ23168 (FLJ23168), mRNA
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NM_025005	Homo sapiens hypothetical protein FLJ13315 (FLJ13315), mRNA
NM_024998	Homo sapiens hypothetical protein FLJ12704 (FLJ12704), mRNA
NM_024994	Homo sapiens hypothetical protein FLJ12595 (FLJ12595), mRNA
NM_024977	Homo sapiens hypothetical protein FLJ12078 (FLJ12078), mRNA
NM_024976	Homo sapiens hypothetical protein FLJ11996 (FLJ11996), mRNA
NM_024956	Homo sapiens hypothetical protein FLJ23375 (FLJ23375), mRNA
NM_024944	Homo sapiens chromosome 21 open reading frame 68 (C21orf68), mRNA
NM_024942	Homo sapiens hypothetical protein FLJ13490 (FLJ13490), mRNA
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NM_024935	Homo sapiens hypothetical protein FLJ13687 (FLJ13687), mRNA
NM_024920	Homo sapiens hypothetical protein FLJ14281 (FLJ14281), mRNA
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NM_024897	Homo sapiens hypothetical protein FLJ22672 (FLJ22672), mRNA
NM_024889	Homo sapiens hypothetical protein FLJ23537 (FLJ23537), mRNA
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NM_024880	Homo sapiens hypothetical protein FLJ23556 (FLJ23556), mRNA
NM_024864	Homo sapiens hypothetical protein FLJ22578 (FLJ22578), mRNA
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NM_024837	Homo sapiens hypothetical protein FLJ21472 (FLJ21472), mRNA
NM_024835	Homo sapiens C3HC4-type zinc finger protein (LZK1), mRNA
NM_024815	Homo sapiens hypothetical protein FLJ22494 (FLJ22494), mRNA
	

NM_024813	Homo sapiens hypothetical protein FLJ13150 (FLJ13150), mRNA
NM_024811	Homo sapiens hypothetical protein FLJ12529 (FLJ12529), mRNA
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NM_024808	Homo sapiens hypothetical protein FLJ22624 (FLJ22624), mRNA
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NM_024806	Homo sapiens hypothetical protein FLJ23554 (FLJ23554), mRNA
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NM 024746	Homo sapiens hypothetical protein FLJ13840 (FLJ13840), mRNA
NM 024732	Homo sapiens hypothetical protein FLJ14351 (FLJ14351), mRNA
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NM 024727	Homo sapiens hypothetical protein FLJ23259 (FLJ23259), mRNA
NM 024722	Homo sapiens hypothetical protein FLJ13322 (FLJ13322), mRNA
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NM 024715	Homo sapiens hypothetical protein FLJ22625 (FLJ22625), mRNA
NM 024709	Homo sapiens hypothetical protein FLJ14146 (FLJ14146), mRNA
NM 024705	Homo sapiens hypothetical protein FLJ13639 (FLJ13639), mRNA
NM 024703	Homo sapiens hypothetical protein FLJ22593 (FLJ22593), mRNA
NM 024701	Homo sapiens ankyrin repeat and SOCS box-containing 13 (ASB13), mRNA
NM 024700	Homo sapiens Smad nuclear interacting protein (SNIP1), mRNA
NM 024695	Homo sapiens hypothetical protein FLJ13993 (FLJ13993), mRNA
NM 024693	Homo sapiens hypothetical protein FLJ20909 (FLJ20909), mRNA
NM 024688	Homo sapiens hypothetical protein FLJ13031 (FLJ13031), mRNA
NM 024686	Homo sapiens hypothetical protein FLJ23033 (FLJ23033), mRNA
NM 024678	Homo sapiens hypothetical protein FLJ23441 (FLJ23441), mRNA
NM 024675	Homo sapiens hypothetical protein FLJ21816 (FLJ21816), mRNA
NM 024672	Homo sapiens hypothetical protein FLJ23320 (FLJ23320), mRNA
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NM 024654	Homo sapiens hypothetical protein FLJ23323 (FLJ23323), mRNA
NM 024650	Homo sapiens hypothetical protein FLJ22531 (FLJ22531), mRNA
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NM 024647	Homo sapiens hypothetical protein FLJ13287 (FLJ13287), mRNA
NM 024640	Homo sapiens hypothetical protein FLJ23476 (FLJ23476), mRNA
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	related protein (FLJ23153), mRNA
NM 024628	Homo sapiens hypothetical protein FLJ23188 (FLJ23188), mRNA
NM 024627	Homo sapiens hypothetical protein FLJ21125 (FLJ21125), mRNA
NM_024626	Homo sapiens hypothetical protein FLJ22418 (FLJ22418), mRNA
NM 024624	Homo sapiens hypothetical protein FLJ22116 (FLJ22116), mRNA
NM_024616	Homo sapiens hypothetical protein FLJ23186 (FLJ23186), mRNA
NM 024615	Homo sapiens hypothetical protein FLJ21308 (FLJ21308), mRNA
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NM_024613	Homo sapiens phafin 2 (FLJ13187), mRNA
NM_024610	Homo sapiens hypothetical protein FLJ22623 (FLJ22623), mRNA
NM_024609	Homo sapiens hypothetical protein FLJ21841 (FLJ21841), mRNA
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NM_024605	Homo sapiens hypothetical protein FLJ20896 (FLJ20896), mRNA
NM_024602	Homo sapiens hypothetical protein FLJ21156 (FLJ21156), mRNA
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NM_024507	Homo sapiens hypothetical protein MGC10791 (MGC10791), mRNA
NM_015288	Homo sapiens KIAA0239 protein (KIAA0239), mRNA
NM_024419	Homo sapiens Phosphatidylglycerophosphate Synthase (PGS1), mRNA
NM_024345	Homo sapiens hypothetical protein MGC10765 (MGC10765), mRNA
NM_024340	Homo sapiens hypothetical protein MGC4179 (MGC4179), mRNA
NM_024330	Homo sapiens hypothetical protein MGC4365 (MGC4365), mRNA
NM_024326	Homo sapiens hypothetical protein MGC11279 (MGC11279), mRNA
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NM_024295	Homo sapiens hypothetical protein MGC3067 (MGC3067), mRNA
NM_020062	Homo sapiens SLC2A4 regulator (SLC2A4RG), mRNA
NM_018491	Homo sapiens COBW-like protein (LOC55871), mRNA
NM_024116	Homo sapiens hypothetical protein MGC5306 (MGC5306), mRNA
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NM_024113	Homo sapiens hypothetical protein MGC4707 (MGC4707), mRNA
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NM_024084	Homo sapiens hypothetical protein MGC3196 (MGC3196), mRNA
NM_024072 NM_024067	Homo sapiens hypothetical protein MGC2835 (MGC2835), mRNA
NM 024067	Homo sapiens hypothetical protein MGC2718 (MGC2718), mRNA
NM 024040	Homo sapiens hypothetical protein MGC5347 (MGC5347), mRNA
NM 024036	Homo sapiens hypothetical protein MGC2491 (MGC2491), mRNA
NM_024036 NM_015450	Homo sapiens hypothetical protein MGC3103 (MGC3103), mRNA
NM_021249	Homo sapiens protection of telomeres 1 (POT1), mRNA
NM_023932	Homo sapiens sorting nexin 6 (SNX6), mRNA Homo sapiens hypothetical protein MCC2487 04CC2487
NM_023930	Homo sapiens hypothetical protein MGC2487 (MGC2487), mRNA
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NM_014045	Homo sapiens DKFZP564C1940 protein (DKFZP564C1940), mRNA
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NM_023923	Homo sapiens hypothetical protein FLJ13171 (FLJ13171), mRNA
NM_019054	Homo sapiens hypothetical protein MGC5560 (MGC5560), mRNA
NM_023070	Homo sapiens hypothetical protein (LOC65243), mRNA
NM_023015	Homo sapiens hypothetical protein FLJ21919 (FLJ21919), mRNA
NM_022899	Homo sapiens likely ortholog of mouse actin-related protein 8 homolog (S.
	cerevisiae) (FLJ12934), mRNA
NM_022836	Homo sapiens DNA cross-link repair 1B (PSO2 homolog, S. cerevisiae)
	(DCLRE1B), mRNA
NM_022831	Homo sapiens hypothetical protein FLJ12806 (FLJ12806), mRNA
NM_022828	Homo sapiens hypothetical protein FLJ21940 (FLJ21940), mRNA
NM_022822	Homo sapiens hypothetical protein FLJ12387 similar to kinesin light chain (FLJ12387), mRNA
NM_022784	Homo sapiens hypothetical protein FLJ12476 (FLJ12476), mRNA
NM_022783	Homo sapiens hypothetical protein FLJ12428 (FLJ12428), mRNA
NM_022774	Homo sapiens hypothetical protein FLJ21144 (FLJ21144), mRNA
NM_022765	Homo sapiens hypothetical protein FLJ11937 (FLJ11937), mRNA
NM_022764	Homo sapiens hypothetical protein FLJ12998 (FLJ12998), mRNA
NM_022758	Homo sapiens hypothetical protein FLJ22195 (FLJ22195), mRNA
NM 022753	Homo sapiens hypothetical protein FLJ12903 (FLJ12903), mRNA
NM 022749	Homo sapiens retinoic acid induced 16 (RAI16), mRNA
NM 022746	Homo sapiens hypothetical protein FLJ22390 (FLJ22390), mRNA
NM 022728	Homo sapiens neurogenic differentiation 6 (NEUROD6), mRNA
NM 022496	Homo sapiens hypothetical protein FLJ13433 (FLJ13433), mRNA
NM_022490	Homo sapiens hypothetical protein FLJ13390 similar to PAF53 (FLJ13390), mRNA
NM 022484	Homo sapiens hypothetical protein FLJ13576 (FLJ13576), mRNA
NM 022483	Homo sapiens hypothetical protein FLJ21657 (FLJ21657), mRNA
NM 022473	Homo sapiens zinc finger protein 106 (ZFP106), mRNA
NM 022471	Homo sapiens hypothetical protein FLJ13057 similar to germ cell-less
	(FLJ13057), mRNA
NM_022463	Homo sapiens nucleoredoxin 1 (NXN), mRNA
NM_022462	Homo sapiens hypothetical protein FLJ14033 similar to hypoxia inducible factor
	3, alpha subunit (HIF-3A), mRNA
NM_022461	Homo sapiens hypothetical protein FLJ21939 similar to 5-azacytidine induced gene 2 (FLJ21939), mRNA
NM_022453	Homo sapiens ring finger protein 25 (RNF25), mRNA
NM 022374	Homo sapiens likely ortholog of mouse ADP-ribosylation-like factor 6
=	interacting protein 2 (FLJ23293), mRNA
NM 022371	Homo sapiens ATP-dependant interferon responsive (ADIR), mRNA
NM 022369	Homo sapiens hypothetical protein FLJ12541 similar to Stra6 (FLJ12541),
	mRNA
NM_022367	Homo sapiens hypothetical protein FLJ12287 similar to semaphorins (FLJ12287), mRNA
NM 022359	Homo sapiens similar to rat myomegalin (LOC64182), mRNA
NM 022356	Homo sapiens growth suppressor 1 (GROS1), mRNA
NM 022354	Homo sapiens spermatogenesis associated 1 (SPATA1), mRNA
NM 022347	Homo sapiens IFRG15 protein (IFRG15), mRNA
NM_022341	Homo sapiens peptide deformylase-like protein (LOC64146), mRNA
NM 022164	Homo sapiens P3ECSL (LIECG3), mRNA
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NM_022147 Homo sapiens 28kD interferon responsive protein (IFRG28), mRNA NM_022140 Homo sapiens erythrocyte protein band 4.1-like 4 (EPB41L4), mRNA NM_022126 Homo sapiens sorting nexin 16 (SNX16), mRNA NM_022126 Homo sapiens sphospholysine phosphohistidine inorganic pyrophosphate phosphatase (LHPP), mRNA NM_022097 Homo sapiens hepatocellular carcinoma antigen gene 520 (LOC63928), mRNA NM_022094 Homo sapiens hypothetical protein FLJ20871 similar to FSP27 (FLJ20871), mRNA NM_022094 Homo sapiens transposon-derived Buster3 transposase-like (LOC63920), mRNA NM_022074 Homo sapiens hypothetical protein FLJ222794 (FLJ22794), mRNA NM_022074 Homo sapiens hypothetical protein FLJ20967 (FLJ20967), mRNA NM_022063 Homo sapiens hypothetical protein FLJ13188 (FLJ13188), mRNA NM_022064 Homo sapiens hypothetical protein FLJ12816 (FLJ12816), mRNA NM_021034 Homo sapiens estrogen regulated gene 1 (ERG-1), mRNA NM_021945 Homo sapiens hypothetical protein FLJ12174 (FLJ22174), mRNA NM_021944 Homo sapiens hypothetical protein FLJ12154 (FLJ12154), mRNA NM_021941 Homo sapiens hypothetical protein FLJ12154 (FLJ21324), mRNA NM_021928 Homo sapiens hypothetical protein FLJ121324 (FLJ21324), mRNA NM_021927 Homo sapiens hypothetical protein FLJ12180 (FLJ13220), mRNA NM_021927 Homo sapiens hypothetical protein FLJ13220 (FLJ13220), mRNA NM_021825 Homo sapiens hypothetical protein FLJ1320 (FLJ13220), mRNA NM_021639 Homo sapiens hypothetical protein SP192 (SP192), mRNA NM_021637 Homo sapiens hypothetical protein FLJ14084 (FLJ14084), mRNA NM_021639 Homo sapiens hypothetical protein FLJ14084 (FLJ14084), mRNA NM_021631 Homo sapiens potassium intermediate/small conductance calcium-activated channel, subfamily N, member 2 (KCNN2), mRNA NM_021170 Homo sapiens minor histocompatibility antigen HB-1 (HB-1), mRNA NM_021174 Homo sapiens angiopoietin-like factor (CDT6), mRNA
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NM 021927 Homo sapiens hypothetical protein FLJ13220 (FLJ13220), mRNA NM 021925 Homo sapiens hypothetical protein FLJ21820 (FLJ21820), mRNA NM 021825 Homo sapiens hypothetical protein MDS025 (MDS025), mRNA NM 015622 Homo sapiens CGI-43 protein (LOC51622), mRNA NM 021639 Homo sapiens hypothetical protein SP192 (SP192), mRNA NM 021637 Homo sapiens hypothetical protein FLJ14084 (FLJ14084), mRNA NM 021614 Homo sapiens potassium intermediate/small conductance calcium-activated channel, subfamily N, member 2 (KCNN2), mRNA NM 021182 Homo sapiens minor histocompatibility antigen HB-1 (HB-1), mRNA NM 021170 Homo sapiens bHLH factor Hes4 (LOC57801), mRNA
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NM 021825 Homo sapiens hypothetical protein MDS025 (MDS025), mRNA NM 015622 Homo sapiens CGI-43 protein (LOC51622), mRNA NM 021639 Homo sapiens hypothetical protein SP192 (SP192), mRNA NM 021637 Homo sapiens hypothetical protein FLJ14084 (FLJ14084), mRNA NM 021614 Homo sapiens potassium intermediate/small conductance calcium-activated channel, subfamily N, member 2 (KCNN2), mRNA NM 021182 Homo sapiens minor histocompatibility antigen HB-1 (HB-1), mRNA NM 021170 Homo sapiens bHLH factor Hes4 (LOC57801), mRNA
NM 015622 Homo sapiens CGI-43 protein (LOC51622), mRNA NM 021639 Homo sapiens hypothetical protein SP192 (SP192), mRNA NM 021637 Homo sapiens hypothetical protein FLJ14084 (FLJ14084), mRNA NM 021614 Homo sapiens potassium intermediate/small conductance calcium-activated channel, subfamily N, member 2 (KCNN2), mRNA NM 021182 Homo sapiens minor histocompatibility antigen HB-1 (HB-1), mRNA NM 021170 Homo sapiens bHLH factor Hes4 (LOC57801), mRNA
NM 021639 Homo sapiens hypothetical protein SP192 (SP192), mRNA NM 021637 Homo sapiens hypothetical protein FLJ14084 (FLJ14084), mRNA NM 021614 Homo sapiens potassium intermediate/small conductance calcium-activated channel, subfamily N, member 2 (KCNN2), mRNA NM 021182 Homo sapiens minor histocompatibility antigen HB-1 (HB-1), mRNA NM 021170 Homo sapiens bHLH factor Hes4 (LOC57801), mRNA
NM 021637 Homo sapiens hypothetical protein FLJ14084 (FLJ14084), mRNA NM_021614 Homo sapiens potassium intermediate/small conductance calcium-activated channel, subfamily N, member 2 (KCNN2), mRNA NM_021182 Homo sapiens minor histocompatibility antigen HB-1 (HB-1), mRNA NM_021170 Homo sapiens bHLH factor Hes4 (LOC57801), mRNA
NM_021614 Homo sapiens potassium intermediate/small conductance calcium-activated channel, subfamily N, member 2 (KCNN2), mRNA NM_021182 Homo sapiens minor histocompatibility antigen HB-1 (HB-1), mRNA NM_021170 Homo sapiens bHLH factor Hes4 (LOC57801), mRNA
channel, subfamily N, member 2 (KCNN2), mRNA NM_021182 Homo sapiens minor histocompatibility antigen HB-1 (HB-1), mRNA NM_021170 Homo sapiens bHLH factor Hes4 (LOC57801), mRNA
NM_021182 Homo sapiens minor histocompatibility antigen HB-1 (HB-1), mRNA NM_021170 Homo sapiens bHLH factor Hes4 (LOC57801), mRNA
NM_021170 Homo sapiens bHLH factor Hes4 (LOC57801), mRNA
NM_021146 Homo sapiens angiopoietin-like factor (CDT6), mRNA
NM_005146 Homo sapiens squamous cell carcinoma antigen recognised by T cells (SART1),
mRNA
NM_021079 Homo sapiens N-myristoyltransferase 1 (NMT1), mRNA
NM_021046 Homo sapiens UHS KerB (LOC57830), mRNA
NM_021018 Homo sapiens H3 histone family, member I (H3FI), mRNA
NM_006643 Homo sapiens serologically defined colon cancer antigen 3 (SDCCAG3), mRNA
NM_017569 Homo sapiens transcription factor (p38 interacting protein) (P38IP), mRNA
NM_015239 Homo sapiens KIAA1035 protein (KIAA1035), mRNA
NM_014977 Homo sapiens KIAA0670 protein/acinus (KIAA0670), mRNA
NM_015176 Homo sapiens KIAA0483 protein (KIAA0483), mRNA
NM_014610 Homo sapiens KIAA0088 protein (KIAA0088), mRNA
NM_015516 Homo sapiens hypothetical protein, estradiol-induced (E2IG4), mRNA
NM_015388 Homo sapiens DKFZP566C243 protein (DKFZP566C243), mRNA
NM_015679 Homo sapiens hypothetical protein (CLONE24922), mRNA
NM_014409 Homo sapiens TAF5-like RNA polymerase II, p300/CBP-associated factor
(PCAF)-associated factor, 65 kD (TAF5L), mRNA
NM_014368 Homo sapiens LIM homeobox protein 6 (LHX6), mRNA
NM_014315 Homo sapiens host cell factor homolog (LCP), mRNA
NM_012414 Homo sapiens rab3 GTPase-activating protein, non-catalytic subunit (150kD)
(RAB3-GAP150), mRNA
NM_012219 Homo sapiens muscle RAS oncogene homolog (MRAS), mRNA
NM_007375 Homo sapiens TAR DNA binding protein (TARDBP), mRNA

NM_007074	Homo sapiens coronin, actin binding protein, 1A (CORO1A), mRNA
NM_006927	Homo sapiens sialyltransferase 4B (beta-galactosidase alpha-2,3-
	sialytransferase) (SIAT4B), mRNA
NM_006861	Homo sapiens RAB35, member RAS oncogene family (RAB35), mRNA
NM_006502	Homo sapiens polymerase (DNA directed), eta (POLH), mRNA
NM_005710	Homo sapiens polyglutamine binding protein 1 (PQBP1), mRNA
NM_005168	Homo sapiens ras homolog gene family, member E (ARHE), mRNA
NM_004190	Homo sapiens lipase, gastric (LIPF), mRNA
NM_004132	Homo sapiens hyaluronan binding protein 2 (HABP2), mRNA
NM_004492	Homo sapiens general transcription factor IIA, 2 (12kD subunit) (GTF2A2), mRNA
NM_004824	Homo sapiens chromodomain protein, Y chromosome-like (CDYL), mRNA
NM_003969	Homo sapiens ubiquitin-conjugating enzyme E2M (UBC12 homolog, yeast) (UBE2M), mRNA
NM_002711	Homo sapiens protein phosphatase 1, regulatory (inhibitor) subunit 3A (glycogen and sarcoplasmic reticulum binding subunit, skeletal muscle) (PPP1R3A), mRNA
NM_003847	Homo sapiens peroxisomal biogenesis factor 11A (PEX11A), mRNA
NM_002004	Homo sapiens farnesyl diphosphate synthase (farnesyl pyrophosphate synthetase, dimethylallyltranstransferase, geranyltranstransferase) (FDPS), mRNA
NM_019111	Homo sapiens major histocompatibility complex, class II, DR alpha (HLA-DRA), mRNA
NM_002120	Homo sapiens major histocompatibility complex, class II, DO beta (HLA-DOB), mRNA
NM_002118	Homo sapiens major histocompatibility complex, class II, DM beta (HLA-DMB), mRNA
NM_002125	Homo sapiens major histocompatibility complex, class II, DR beta 5 (HLA-DRB5), mRNA
NM_021983	Homo sapiens major histocompatibility complex, class II, DR beta 4 (HLA-DRB4), mRNA
NM_022555	Homo sapiens major histocompatibility complex, class II, DR beta 3 (HLA-DRB3), mRNA
NM_005962	Homo sapiens MAX interacting protein 1 (MXI1), transcript variant 1, mRNA
NM_130439	Homo sapiens MAX interacting protein 1 (MXI1), transcript variant 2, mRNA
NM_080923	Homo sapiens protein tyrosine phosphatase, receptor type, C (PTPRC), transcript variant 4, mRNA
NM_080922	Homo sapiens protein tyrosine phosphatase, receptor type, C (PTPRC), transcript variant 3, mRNA
NM_080921	Homo sapiens protein tyrosine phosphatase, receptor type, C (PTPRC), transcript variant 2, mRNA
NM_130386	Homo sapiens collectin sub-family member 12 (COLEC12), transcript variant I, mRNA
NM_030781	Homo sapiens collectin sub-family member 12 (COLEC12), transcript variant II, mRNA
NM_130778	Homo sapiens collagen, type XVII, alpha 1 (COL17A1), transcript variant short, mRNA
NM_000494	Homo sapiens collagen, type XVII, alpha 1 (COL17A1), transcript variant long, mRNA
NM 001856	Homo sapiens collagen, type XVI, alpha 1 (COL16A1), mRNA
NM 001855	Homo sapiens collagen, type XVI, alpha 1 (COL10A1), mRNA
NM_058166	Homo sapiens tripartite motif-containing 6 (TRIM6), mRNA
NM_002838	Homo sapiens protein tyrosine phosphatase, receptor type, C (PTPRC), transcript
	ranscript

	variant 1, mRNA
NB4 120200	
NM_130390	Homo sapiens tripartite motif-containing 34 (TRIM34), transcript variant 3,
3B 6 100000	mRNA
NM_130389	Homo sapiens tripartite motif-containing 34 (TRIM34), transcript variant 2,
277.6 001616	mRNA
NM_021616	Homo sapiens tripartite motif-containing 34 (TRIM34), transcript variant 1,
ND 6 000050	mRNA
NM_030950	Homo sapiens ret finger protein (RFP), transcript variant beta, mRNA
NM_130785	Homo sapiens TPTE and PTEN homologous inositol lipid phosphatase (TPIP),
377	mRNA
NM_130784	Homo sapiens hypothetical gene supported by AY027807; AY027808
L	(LOC93426), mRNA
NM_130783	Homo sapiens similar to neuronal tetraspanin (LOC90139), mRNA
NM_130782	Homo sapiens regulator of G-protein signalling 18 (RGS18), mRNA
NM_130781	Homo sapiens (RAB24), mRNA
NM_130772	Homo sapiens S100Z protein (S100Z), mRNA
NM_130769	Homo sapiens glycoprotein alpha 2 (GPA2), mRNA
NM_130770	Homo sapiens 5-hydroxytryptamine receptor 3 subunit C (HTR3C), mRNA
NM_130768	Homo sapiens GASZ (GASZ), mRNA
NM_130767	Homo sapiens cytosolic acetyl-CoA hydrolase (CACH-1), mRNA
NM_130773	Homo sapiens caspr5 protein (caspr5), mRNA
NM_006510	Homo sapiens ret finger protein (RFP), transcript variant alpha, mRNA
NM_033554	Homo sapiens major histocompatibility complex, class II, DP alpha 1 (HLA-
	DPA1), mRNA
NM_033282	Homo sapiens opsin 4 (melanopsin) (OPN4), mRNA
NM_032035	Homo sapiens MSTP031 protein (MSTP031), mRNA
NM_017882	Homo sapiens ceroid-lipofuscinosis, neuronal 6, late infantile, variant (CLN6),
	<u> </u> mRNA
NM_006983	Homo sapiens matrix metalloproteinase 23B (MMP23B), mRNA
NM_005608	Homo sapiens protein tyrosine phosphatase, receptor type, C-associated protein
	(PTPRCAP), mRNA
NM_004659	Homo sapiens matrix metalloproteinase 23A (MMP23A), mRNA
NM_025091	Homo sapiens hypothetical protein FLJ13330 (FLJ13330), mRNA
NM_130759	Homo sapiens immunity associated protein 1 (IMAP1), mRNA
NM_019841	Homo sapiens transient receptor potential cation channel, subfamily V, member
	5 (TRPV5), mRNA
NM_017584	Homo sapiens aldehyde reductase (aldose reductase) like 6 (ALDRL6), mRNA
NM_017436	Homo sapiens alpha 1,4-galactosyltransferase (A4GALT), mRNA
NM_006480	Homo sapiens regulator of G-protein signalling 14 (RGS14), mRNA
NM_013357	Homo sapiens purine-rich element binding protein G (PURG), mRNA
NM_016155	Homo sapiens matrix metalloproteinase 17 (membrane-inserted) (MMP17).
37.6.000000	mRNA
NM_002813	Homo sapiens proteasome (prosome, macropain) 26S subunit, non-ATPase, 9
37.6.001.5.0	(PSMD9), mRNA
NM_024549	Homo sapiens hypothetical protein FLJ21127 (FLJ21127), mRNA
NM_130441	Homo sapiens dendritic cell lectin b (DLEC), mRNA
NM_015409	Homo sapiens E1A binding protein p400 (EP400), mRNA
NM_003702	Homo sapiens regulator of G-protein signalling 20 (RGS20), mRNA
NM_016113	Homo sapiens transient receptor potential cation channel, subfamily V, member
	2 (TRPV2), mRNA
NM_015530	Homo sapiens likely ortholog of rat golgi stacking protein homolog GRASP55
	(GKASP55), mRNA
	(GRASP55), mRNA

ND4 005072	Homo sapiens regulator of G-protein signalling 19 (RGS19), mRNA
NM_005873	
NM_130469	Homo sapiens Jun dimerization protein 2 (jdp2), mRNA
NM_130468	Homo sapiens dermatan-4-sulfotransferase-1 (D4ST-1), mRNA
NM_130467	Homo sapiens PAGE-5 protein (PAGE-5), mRNA
NM_130463	Homo sapiens ATPase, H+ transporting, lysosomal (vacuolar proton pump) (ATP6G), mRNA
NM_130459	Homo sapiens torsin family 2, member A (TOR2A), mRNA
NM_021070	Homo sapiens latent transforming growth factor beta binding protein 3 (LTBP3), mRNA
NM_020865	Homo sapiens DEAD/H (Asp-Glu-Ala-Asp/His) box polypeptide 36 (DDX36), mRNA
NM_016304	Homo sapiens 60S ribosomal protein L30 isolog (LOC51187), mRNA
NM_130443	Homo sapiens dipeptidylpeptidase III (DPP3), transcript variant 2, mRNA
NM_005700	Homo sapiens dipeptidylpeptidase III (DPP3), transcript variant 1, mRNA
NM 018152	Homo sapiens chromosome 20 open reading frame 12 (C20orf12), mRNA
NM 006027	Homo sapiens exonuclease 1 (EXO1), transcript variant 1, mRNA
NM_003686	Homo sapiens exonuclease 1 (EXO1), transcript variant 3, mRNA
NM_130398	Homo sapiens exonuclease 1 (EXO1), transcript variant 2, mRNA
NM_002837	Homo sapiens protein tyrosine phosphatase, receptor type, B (PTPRB), mRNA
NM_000775	Homo sapiens cytochrome P450, subfamily IIJ (arachidonic acid epoxygenase) polypeptide 2 (CYP2J2), mRNA
NM_053056	Homo sapiens cyclin D1 (PRAD1 parathyroid adenomatosis 1) (CCND1), mRNA
NM_012090	Homo sapiens microtubule-actin crosslinking factor 1 (MACF1), transcript variant 1, mRNA
NM 017625	Homo sapiens intelectin (ITLN), mRNA
NM_015839	Homo sapiens ficolin (collagen/fibrinogen domain containing lectin) 2 (hucolin) (FCN2), transcript variant SV3, mRNA
NM_015838	Homo sapiens ficolin (collagen/fibrinogen domain containing lectin) 2 (hucolin) (FCN2), transcript variant SV2, mRNA
NM_015837	Homo sapiens ficolin (collagen/fibrinogen domain containing lectin) 2 (hucolin) (FCN2), transcript variant SV1, mRNA
NM 002003	Homo sapiens ficolin (collagen/fibrinogen domain containing) 1 (FCN1), mRNA
NM 016327	Homo sapiens ureidopropionase, beta (UPB1), mRNA
NM_016328	Homo sapiens GTF2I repeat domain containing 1 (GTF2IRD1), transcript variant 1, mRNA
NM_004108	Homo sapiens ficolin (collagen/fibrinogen domain containing lectin) 2 (hucolin) (FCN2), transcript variant SV0, mRNA
NM 002318	Homo sapiens lysyl oxidase-like 2 (LOXL2), mRNA
NM_130396	Homo sapiens WNT1 inducible signaling pathway protein 3 (WISP3), transcript variant 2, mRNA
NM_003880	Homo sapiens WNT1 inducible signaling pathway protein 3 (WISP3), transcript variant 1, mRNA
NM 003881	Homo sapiens WNT1 inducible signaling pathway protein 2 (WISP2), mRNA
NM_080838	Homo sapiens WNT1 inducible signaling pathway protein 1 (WISP1), transcript variant 2, mRNA
NM_003882	Homo sapiens WNT1 inducible signaling pathway protein 1 (WISP1), transcript variant 1, mRNA
NM_000651	Homo sapiens complement component (3b/4b) receptor 1, including Knops blood group system (CR1), transcript variant S, mRNA
NM_000573	Homo sapiens complement component (3b/4b) receptor 1, including Knops blood group system (CR1), transcript variant F, mRNA

NM_006069	Homo sapiens murine retrovirus integration site 1 homolog (MRVII), transcript variant 1, mRNA
NM_130385	Homo sapiens murine retrovirus integration site 1 homolog (MRVII), transcript variant 2, mRNA
NM 018492	Homo sapiens T-LAK cell-originated protein kinase (TOPK), mRNA
NM 002462	Homo sapiens myxovirus (influenza virus) resistance 1, interferon-inducible
_	protein p78 (mouse) (MX1), mRNA
NM 015920	Homo sapiens ribosomal protein S27-like (RPS27L), mRNA
NM 016183	Homo sapiens ribosomal protein, large, P0-like (RPLP0L), mRNA
NM 080746	Homo sapiens ribosomal protein L10-like (RPL10L), mRNA
NM_032236	Homo sapiens FLJ23277 protein (FLJ23277), mRNA
NM_032784	Homo sapiens thrombospondin (FLJ14440), mRNA
NM_080731	Homo sapiens intermediate filament-like MGC:2625 (DKFZP586I2223).
	transcript variant 3, mRNA
NM_080730	Homo sapiens intermediate filament-like MGC:2625 (DKFZP586I2223),
	transcript variant 2, mRNA
NM_015945	Homo sapiens ovarian cancer overexpressed 1 (OVCOV1), mRNA
NM_018018	Homo sapiens solute carrier family 38, member 4 (SLC38A4), mRNA
NM_022451	Homo sapiens AD24 protein (AD24), mRNA
NM_020830	Homo sapiens phosphoinositide-binding protein SR1 (FENS-1), mRNA
NM_033630	Homo sapiens SCAN domain containing 1 (SCAND1), transcript variant 2, mRNA
NM_016558	Homo sapiens SCAN domain containing 1 (SCAND1), transcript variant 1, mRNA
NM_015438	Homo sapiens intermediate filament-like MGC:2625 (DKFZP586I2223), transcript variant 1, mRNA
NM_007371	Homo sapiens bromodomain containing 3 (BRD3), mRNA
NM 005104	Homo sapiens bromodomain containing 2 (BRD2), mRNA
NM_005031	Homo sapiens FXYD domain containing ion transport regulator 1 (phospholemman) (FXYD1), transcript variant a, mRNA
NM_021902	Homo sapiens FXYD domain containing ion transport regulator 1
1414_021302	(phospholemman) (FXYD1), transcript variant b, mRNA
NM_014164	Homo sapiens FXYD domain-containing ion transport regulator 5 (FXYD5),
NM_002463	mRNA
NM 014577	Homo sapiens myxovirus (influenza virus) resistance 2 (mouse) (MX2), mRNA
NM 021004	Homo sapiens bromodomain containing 1 (BRD1), mRNA
_	Homo sapiens peroxisomal short-chain alcohol dehydrogenase (humNRDR), mRNA
NM_020399	Homo sapiens PDZ/coiled-coil domain binding partner for the rho-family GTPase TC10 (PIST), mRNA
NM_017935	Homo sapiens hypothetical protein FLJ20706 (BANK), mRNA
NM_018244	Homo sapiens chromosome 20 open reading frame 44 (C20orf44), mRNA
NM_016100	Homo sapiens N-acetyltransferase 5 (ARD1 homolog, S. cerevisiae) (NAT5), mRNA
NM_016045	Homo sapiens chromosome 20 open reading frame 45 (C20orf45), mRNA
NM_007363	Homo sapiens non-POU domain containing, octamer-binding (NONO), mRNA
NM_002438	Homo sapiens mannose receptor, C type 1 (MRC1), mRNA
NM_015092	Homo sapiens PI-3-kinase-related kinase SMG-1 (SMG1), mRNA
NM_018993	Homo sapiens RAB5 interacting protein 2 (RIN2), mRNA
NM_080841	Homo sapiens protein tyrosine phosphatase, receptor type, A (PTPRA), transcript variant 3, mRNA
NM_080840	Homo sapiens protein tyrosine phosphatase, receptor type, A (PTPRA),
	, , , , , , , , , , , , , , , , , , ,

	transcript variant 2, mRNA
NM_002836	Homo sapiens protein tyrosine phosphatase, receptor type, A (PTPRA),
	transcript variant 1, mRNA
NM 024832	Homo sapiens RAB5 interacting protein 3 (RIN3), mRNA
NM 023915	Homo sapiens G protein-coupled receptor 87 (GPR87), mRNA
NM 003029	Homo sapiens SHC (Src homology 2 domain containing) transforming protein 1
_	(SHC1), mRNA
NM 018490	Homo sapiens G protein-coupled receptor 48 (GPR48), mRNA
NM 016020	Homo sapiens homolog of yeast mitochondrial transcription factor B (mtTFB),
	mRNA
NM 014475	Homo sapiens dihydrodiol dehydrogenase (dimeric) (DHDH), mRNA
NM 006065	Homo sapiens signal-regulatory protein beta 1 (SIRPB1), mRNA
NM 005527	Homo sapiens heat shock 70kD protein 1-like (HSPA1L), mRNA
NM_004648	Homo sapiens protein tyrosine phosphatase, non-receptor type substrate 1
	(PTPNS1), mRNA
NM_004480	Homo sapiens fucosyltransferase 8 (alpha (1,6) fucosyltransferase) (FUT8),
_	mRNA
NM 003667	Homo sapiens G protein-coupled receptor 49 (GPR49), mRNA
NM 130434	Homo sapiens dipeptidylpeptidase 8 (DPP8), transcript variant 1, mRNA
NM 017743	Homo sapiens dipeptidylpeptidase 8 (DPP8), transcript variant 2, mRNA
NM 002122	Homo sapiens major histocompatibility complex, class II, DQ alpha 1 (HLA-
_	DQA1), mRNA
NM 006442	Homo sapiens DR1-associated protein 1 (negative cofactor 2 alpha) (DRAP1),
-	mRNA
NM_080918	Homo sapiens deoxyguanosine kinase (DGUOK), transcript variant 2, nuclear
_	gene encoding mitochondrial protein, mRNA
NM_080917	Homo sapiens deoxyguanosine kinase (DGUOK), transcript variant 3, nuclear
_	gene encoding mitochondrial protein, mRNA
NM_080916	Homo sapiens deoxyguanosine kinase (DGUOK), transcript variant 1, nuclear
	gene encoding mitochondrial protein, mRNA
NM_080915	Homo sapiens deoxyguanosine kinase (DGUOK), transcript variant 5, nuclear
	gene encoding mitochondrial protein, mRNA
NM_001929	Homo sapiens deoxyguanosine kinase (DGUOK), transcript variant 4, nuclear
	gene encoding mitochondrial protein, mRNA
NM_080815	Homo sapiens collagen, type XIII, alpha 1 (COL13A1), transcript variant 19,
	mRNA
NM_080814	Homo sapiens collagen, type XIII, alpha 1 (COL13A1), transcript variant 18,
	mRNA
NM_080813	Homo sapiens collagen, type XIII, alpha 1 (COL13A1), transcript variant 17,
77.6 000015	mRNA
NM_080812	Homo sapiens collagen, type XIII, alpha 1 (COL13A1), transcript variant 16,
ND 6 0000011	mRNA
NM_080811	Homo sapiens collagen, type XIII, alpha 1 (COL13A1), transcript variant 15,
NTM (000010	mRNA
NM_080810	Homo sapiens collagen, type XIII, alpha 1 (COL13A1), transcript variant 14,
NTM 000000	mRNA
NM_080809	Homo sapiens collagen, type XIII, alpha 1 (COL13A1), transcript variant 13,
NIM OGOGOG	mRNA Home serious cells on the VIII also 1 (COV 12A1) to 12
NM_080808	Homo sapiens collagen, type XIII, alpha 1 (COL13A1), transcript variant 12,
NIM 000007	mRNA Homo seniene collegen to a VIII slabe 1 (COV 12A1) transmitted in 1.11
NM_080807	Homo sapiens collagen, type XIII, alpha 1 (COL13A1), transcript variant 11, mRNA
L	IIIIIII

NM_080806	Homo sapiens collagen, type XIII, alpha 1 (COL13A1), transcript variant 10, mRNA
NM_080805	Homo sapiens collagen, type XIII, alpha 1 (COL13A1), transcript variant 9, mRNA
NM_080804	Homo sapiens collagen, type XIII, alpha 1 (COL13A1), transcript variant 8, mRNA
NM_080803	Homo sapiens collagen, type XIII, alpha 1 (COL13A1), transcript variant 7, mRNA
NM_080802	Homo sapiens collagen, type XIII, alpha 1 (COL13A1), transcript variant 6, mRNA
NM_080801	Homo sapiens collagen, type XIII, alpha 1 (COL13A1), transcript variant 5, mRNA
NM_080800	Homo sapiens collagen, type XIII, alpha 1 (COL13A1), transcript variant 4, mRNA
NM_080799	Homo sapiens collagen, type XIII, alpha 1 (COL13A1), transcript variant 3, mRNA
NM_080798	Homo sapiens collagen, type XIII, alpha 1 (COL13A1), transcript variant 2, mRNA
NM_005203	Homo sapiens collagen, type XIII, alpha 1 (COL13A1), transcript variant 1, mRNA
NM_004395	Homo sapiens drebrin 1 (DBN1), transcript variant 1, mRNA
NM 080881	Homo sapiens drebrin 1 (DBN1), transcript variant 2, mRNA
NM_080792	Homo sapiens brain-immunoglobulin-like molecule with tyrosine-based activation motifs (BIT), mRNA
NM_080816	Homo sapiens signal-regulatory protein beta 2 (SIRPB2), transcript variant 2, mRNA
NM_018556	Homo sapiens signal-regulatory protein beta 2 (SIRPB2), transcript variant 1, mRNA
NM_000787	Homo sapiens dopamine beta-hydroxylase (dopamine beta-monooxygenase) (DBH), mRNA
NM 080426	Homo sapiens GNAS complex locus (GNAS), transcript variant 2, mRNA
NM 080425	Homo sapiens GNAS complex locus (GNAS), transcript variant 3, mRNA
NM 000516	Homo sapiens GNAS complex locus (GNAS), transcript variant 1, mRNA
NM 006571	Homo sapiens novel RGD-containing protein (WS-3), mRNA
NM_080926	Homo sapiens hypothetical protein similar to KIAA0187 gene product (LOC96610), mRNA
NM_080924	Homo sapiens hypothetical protein similar to CGI-67 protein (LOC91219), mRNA
NM_080925	Homo sapiens hypothetical protein similar to topoisomerase (DNA) III beta (H. sapiens) (LOC129020), mRNA
NM_080914	Homo sapiens asialoglycoprotein receptor 2 (ASGR2), transcript variant 3, mRNA
NM_080913	Homo sapiens asialoglycoprotein receptor 2 (ASGR2), transcript variant 2, mRNA
NM_080912	Homo sapiens asialoglycoprotein receptor 2 (ASGR2), transcript variant H2', mRNA
NM_001181	Homo sapiens asialoglycoprotein receptor 2 (ASGR2), transcript variant 1, mRNA
NM_001671	Homo sapiens asialoglycoprotein receptor 1 (ASGR1), mRNA
NM_005065	Homo sapiens sel-1 suppressor of lin-12-like (C. elegans) (SEL1L), mRNA
NM_014978	Homo sapiens VPS10 domain receptor protein SORCS 3 (SORCS3), mRNA
NM 015230	Homo sapiens centaurin, delta 1 (CENTD1), mRNA

NM 052868	Homo sapiens immunoglobulin superfamily, member 8 (IGSF8), mRNA
NM 032782	Homo sapiens hypothetical protein FLJ14428 (TIM3), mRNA
NM 032309	Homo sapiens chromosome 2 open reading frame 9 (C2orf9), mRNA
NM 021625	Homo sapiens transient receptor potential cation channel, subfamily V, member
	4 (TRPV4), mRNA
NM_020960	Homo sapiens G protein-coupled receptor 107 (GPR107), mRNA
NM_024503	Homo sapiens human immunodeficiency virus type I enhancer binding protein 3
	(HIVEP3), mRNA
NM_024112	Homo sapiens chromosome 9 open reading frame 16 (C9orf16), mRNA
NM_015192	Homo sapiens phospholipase C, beta 1 (phosphoinositide-specific) (PLCB1), mRNA
NM 022481	Homo sapiens ARF-GAP, RHO-GAP, ankyrin repeat and plekstrin homology
	domains-containing protein 3 (ARAP3), mRNA
NM 021634	Homo sapiens leucine-rich repeat-containing G protein-coupled receptor 7
	(LGR7), mRNA
NM_013305	Homo sapiens sialyltransferase 8E (alpha-2, 8-polysialytransferase) (SIAT8E), mRNA
NM 019069	Homo sapiens WD repeat domain 5B (WDR5B), mRNA
NM_016179	Homo sapiens transient receptor potential cation channel, subfamily C, member
	4 (TRPC4), mRNA
NM_016592	Homo sapiens GNAS complex locus (GNAS), transcript variant 4, mRNA
NM_014007	Homo sapiens zinc finger protein 297B (ZNF297B), mRNA
NM_012471	Homo sapiens transient receptor potential cation channel, subfamily C, member
	5 (TRPC5), mRNA
NM_012459	Homo sapiens translocase of inner mitochondrial membrane 8 homolog B (yeast) (TIMM8B), mRNA
NM_004621	Homo sapiens transient receptor potential cation channel, subfamily C, member 6 (TRPC6), mRNA
NM_003304	Homo sapiens transient receptor potential cation channel, subfamily C, member 1 (TRPC1), mRNA
NM_002124	Homo sapiens major histocompatibility complex, class II, DR beta 1 (HLA-
	DRB1), mRNA
NM_000972	Homo sapiens ribosomal protein L7a (RPL7A), mRNA
NM_130384	Homo sapiens three prime repair exonuclease 1 (TREX1), transcript variant 6, mRNA
NM_033627	Homo sapiens three prime repair exonuclease 1 (TREX1), transcript variant 2, mRNA
NM_032166	Homo sapiens three prime repair exonuclease 1 (TREX1), transcript variant 5, mRNA
NM 024996	Homo sapiens mitochondrial elongation factor G (EFG1), mRNA
NM_033629	Homo sapiens three prime repair exonuclease 1 (TREX1), transcript variant 4, mRNA
NM_033628	Homo sapiens three prime repair exonuclease 1 (TREX1), transcript variant 3, mRNA
NM_016381	Homo sapiens three prime repair exonuclease 1 (TREX1), transcript variant 1,
NM 031892	
	
NM 013998	Homo sapiens tachykinin, precursor 1 (substance K, substance P, neurokinin 1,
NM 031892 NM 003960 NM 021093 NM 021092 NM 021190	Homo sapiens three prime repair exonuclease 1 (TREX1), transcript variant 1, mRNA Homo sapiens SH3-domain kinase binding protein 1 (SH3KBP1), mRNA Homo sapiens N-acetyltransferase 8 (camello like) (NAT8), mRNA Homo sapiens peptide YY, 2 (seminalplasmin) (PYY2), mRNA Homo sapiens pancreatic polypeptide 2 (PPY2), mRNA Homo sapiens polypyrimidine tract binding protein 2 (PTBP2), mRNA

	neurokinin 2, neuromedin L, neurokinin alpha, neuropeptide K, neuropeptide gamma) (TAC1), transcript variant delta, mRNA
NM 013997	Homo sapiens tachykinin, precursor 1 (substance K, substance P, neurokinin 1,
14141_013997	neurokinin 2, neuromedin L, neurokinin alpha, neuropeptide K, neuropeptide
	gamma) (TAC1), transcript variant gamma, mRNA
NM 013996	Homo sapiens tachykinin, precursor 1 (substance K, substance P, neurokinin 1,
14147_013330	neurokinin 2, neuromedin L, neurokinin alpha, neuropeptide K, neuropeptide
	gamma) (TAC1), transcript variant alpha, mRNA
NM_016235	Homo sapiens G protein-coupled receptor, family C, group 1, member B
	(GPRC5B), mRNA
NM_004630	Homo sapiens splicing factor 1 (SF1), mRNA
NM_000230	Homo sapiens leptin (obesity homolog, mouse) (LEP), mRNA
NM_003185	Homo sapiens TAF4 RNA polymerase II, TATA box binding protein (TBP)-
	associated factor, 135 kD (TAF4), mRNA
NM_003182	Homo sapiens tachykinin, precursor 1 (substance K, substance P, neurokinin 1,
	neurokinin 2, neuromedin L, neurokinin alpha, neuropeptide K, neuropeptide
	gamma) (TAC1), transcript variant beta, mRNA
NM_002772	Homo sapiens protease, serine, 7 (enterokinase) (PRSS7), mRNA
NM_005857	Homo sapiens zinc metalloproteinase (STE24 homolog, yeast) (ZMPSTE24), mRNA
NM_006103	Homo sapiens WAP four-disulfide core domain 2 (WFDC2), transcript variant 1, mRNA
NM_080736	Homo sapiens WAP four-disulfide core domain 2 (WFDC2), transcript variant 2, mRNA
NM 080735	Homo sapiens WAP four-disulfide core domain 2 (WFDC2), transcript variant 5,
	mRNA
NM_080734	Homo sapiens WAP four-disulfide core domain 2 (WFDC2), transcript variant 4, mRNA
NM_080733	Homo sapiens WAP four-disulfide core domain 2 (WFDC2), transcript variant 3, mRNA
NM 021197	Homo sapiens WAP four-disulfide core domain 1 (WFDC1), mRNA
NM 007128	Homo sapiens pre-B lymphocyte gene 1 (VPREB1), mRNA
NM 006373	Homo sapiens vesicle amine transport protein 1 (VATI), mRNA
NM_003105	Homo sapiens sortilin-related receptor, L(DLR class) A repeats-containing (SORL1), mRNA
NM 020777	Homo sapiens VPS10 domain receptor protein (SORCS2), mRNA
NM 052918	Homo sapiens VPS10 domain receptor protein SORCS 1 (SORCS1), mRNA
NM 022553	Homo sapiens SAC2 suppressor of actin mutations 2-like (yeast) (SACM2L),
	transcript variant 2, mRNA
NM 004843	Homo sapiens class I cytokine receptor (WSX1), mRNA
NM 080564	Homo sapiens SAC2 suppressor of actin mutations 2-like (yeast) (SACM2L),
1111_000504	transcript variant 1, mRNA
NM 006711	Homo sapiens RNA binding protein S1, serine-rich domain (RNPS1), transcript
	variant 1, mRNA
NM_080594	Homo sapiens RNA binding protein S1, serine-rich domain (RNPS1), transcript
	variant 2, mRNA
NM 100486	Homo sapiens WW domain-containing adapter with a coiled-coil region (WAC),
	transcript variant 3, mRNA
NM_100264	Homo sapiens WW domain-containing adapter with a coiled-coil region (WAC),
	transcript variant 2, mRNA
NM_016628	Homo sapiens WW domain-containing adapter with a coiled-coil region (WAC),
	transcript variant 1, mRNA

NM_005701	Homo sapiens RNA, U transporter 1 (RNUT1), mRNA
NM_014810	Homo sapiens centrosome-associated protein 350 (CAP350), mRNA
NM_013325	Homo sapiens KIAA0943 protein (Apg4B), mRNA
NM_020235	Homo sapiens bobby sox homolog (Drosophila) (BBX), mRNA
NM_019118	Homo sapiens hypothetical protein RP4-622L5 (RP4-622L5), mRNA
NM_016312	Homo sapiens WW domain binding protein 11 (WBP11), mRNA
NM_018706	Homo sapiens KIAA1630 protein (KIAA1630), mRNA
NM_080599	Homo sapiens regulator of nonsense transcripts 2 (RENT2), transcript variant 1, mRNA
NM_015542	Homo sapiens regulator of nonsense transcripts 2 (RENT2), transcript variant 2, mRNA
NM_002911	Homo sapiens regulator of nonsense transcripts 1 (RENT1), mRNA
NM_002833	Homo sapiens protein tyrosine phosphatase, non-receptor type 9 (PTPN9), mRNA
NM_080589	Homo sapiens protein tyrosine phosphatase, non-receptor type 7 (PTPN7), transcript variant 3, mRNA
NM_080588	Homo sapiens protein tyrosine phosphatase, non-receptor type 7 (PTPN7), transcript variant 2, mRNA
NM_002832	Homo sapiens protein tyrosine phosphatase, non-receptor type 7 (PTPN7), transcript variant 1, mRNA
NM_007039	Homo sapiens protein tyrosine phosphatase, non-receptor type 21 (PTPN21), mRNA
NM_014369	Homo sapiens protein tyrosine phosphatase, non-receptor type 18 (brain-derived) (PTPN18), mRNA
NM_005401	Homo sapiens protein tyrosine phosphatase, non-receptor type 14 (PTPN14), mRNA
NM_002835	Homo sapiens protein tyrosine phosphatase, non-receptor type 12 (PTPN12), mRNA
NM_080685	Homo sapiens protein tyrosine phosphatase, non-receptor type 13 (APO-1/CD95 (Fas)-associated phosphatase) (PTPN13), transcript variant 4, mRNA
NM_080684	Homo sapiens protein tyrosine phosphatase, non-receptor type 13 (APO-1/CD95 (Fas)-associated phosphatase) (PTPN13), transcript variant 3, mRNA
NM_080683	Homo sapiens protein tyrosine phosphatase, non-receptor type 13 (APO-1/CD95 (Fas)-associated phosphatase) (PTPN13), transcript variant 1, mRNA
NM_080601	Homo sapiens protein tyrosine phosphatase, non-receptor type 11 (PTPN11), transcript variant 2, mRNA
NM_002834	Homo sapiens protein tyrosine phosphatase, non-receptor type 11 (PTPN11), transcript variant 1, mRNA
NM_006399	Homo sapiens basic leucine zipper transcription factor, ATF-like (BATF), mRNA
NM_006709	Homo sapiens HLA-B associated transcript 8 (BAT8), transcript variant NG36/G9a, mRNA
NM_033177	Homo sapiens HLA-B associated transcript 4 (BAT4), mRNA
NM_004639	Homo sapiens HLA-B associated transcript 3 (BAT3), transcript variant 1, mRNA
NM_080703	Homo sapiens HLA-B associated transcript 3 (BAT3), transcript variant 3, mRNA
NM_080702	Homo sapiens HLA-B associated transcript 3 (BAT3), transcript variant 2,
\	mRNA
NM_004638	Homo sapiens HLA-B associated transcript 2 (BAT2), transcript variant 2, mRNA

	mRNA
NM_004640	Homo sapiens HLA-B associated transcript 1 (BAT1), transcript variant 1, mRNA
NM_080598	Homo sapiens HLA-B associated transcript 1 (BAT1), transcript variant 2, mRNA
NM_080797	Homo sapiens death associated transcription factor 1 (DATF1), transcript variant 3, mRNA
NM_080796	Homo sapiens death associated transcription factor 1 (DATF1), transcript variant 2, mRNA
NM_022105	Homo sapiens death associated transcription factor 1 (DATF1), transcript variant 1, mRNA
NM 021080	Homo sapiens disabled homolog 1 (Drosophila) (DAB1), mRNA
NM_080760	Homo sapiens dachshund homolog (Drosophila) (DACH), transcript variant 2, mRNA
NM_080759	Homo sapiens dachshund homolog (Drosophila) (DACH), transcript variant 1, mRNA
NM_004392	Homo sapiens dachshund homolog (Drosophila) (DACH), transcript variant 3, mRNA
NM_005996	Homo sapiens T-box 3 (ulnar mammary syndrome) (TBX3), transcript variant 1, mRNA
NM_016569	Homo sapiens T-box 3 (ulnar mammary syndrome) (TBX3), transcript variant 2, mRNA
NM 016954	Homo sapiens T-box 22 (TBX22), mRNA
NM_080701	Homo sapiens three prime repair exonuclease 2 (TREX2), transcript variant 4, mRNA
NM_080700	Homo sapiens three prime repair exonuclease 2 (TREX2), transcript variant 3, mRNA
NM_080699	Homo sapiens three prime repair exonuclease 2 (TREX2), transcript variant 2, mRNA
NM_017518	Homo sapiens three prime repair exonuclease 2 (TREX2), transcript variant 5, mRNA
NM_007205	Homo sapiens three prime repair exonuclease 2 (TREX2), transcript variant 1, mRNA
NM_080632	Homo sapiens similar to yeast Upf3, variant B (UPF3B), transcript variant 1, mRNA
NM_023010	Homo sapiens similar to yeast Upf3, variant B (UPF3B), transcript variant 2, mRNA
NM_080687	Homo sapiens similar to yeast Upf3, variant A (UPF3A), transcript variant 2, mRNA
NM_023011	Homo sapiens similar to yeast Upf3, variant A (UPF3A), transcript variant 1, mRNA
NM_080630	Homo sapiens collagen, type XI, alpha 1 (COL11A1), transcript variant C, mRNA
NM_080629	Homo sapiens collagen, type XI, alpha 1 (COL11A1), transcript variant B, mRNA
NM_001854	Homo sapiens collagen, type XI, alpha 1 (COL11A1), transcript variant A, mRNA
NM_080791	Homo sapiens acid phosphatase, testicular (ACPT), transcript variant A3, mRNA
NM_001639	Homo sapiens amyloid P component, serum (APCS), mRNA
NM_080790	Homo sapiens acid phosphatase, testicular (ACPT), transcript variant A2, mRNA
NM_080789	Homo sapiens acid phosphatase, testicular (ACPT), transcript variant A1, mRNA
NM_033068	Homo sapiens acid phosphatase, testicular (ACPT), transcript variant A, mRNA

NM_001649	Homo sapiens apical protein-like (Xenopus laevis) (APXL), mRNA
NM_014481	Homo sapiens apurinic/apyrimidinic endonuclease-like 2 (APEXL2), nuclear
	gene encoding mitochondrial protein, mRNA
NM_080649	Homo sapiens APEX nuclease (multifunctional DNA repair enzyme) (APEX),
	transcript variant 3, mRNA
NM_080648	Homo sapiens APEX nuclease (multifunctional DNA repair enzyme) (APEX),
	transcript variant 2, mRNA
NM_001641	Homo sapiens APEX nuclease (multifunctional DNA repair enzyme) (APEX),
	transcript variant 1, mRNA
NM_080839	Homo sapiens similar to gamma-glutamyltransferase 1 (LOC91227), mRNA
NM_080927	Homo sapiens endothelial and smooth muscle cell-derived neuropilin-like
	protein (ESDN), mRNA
NM_030969	Homo sapiens hypothetical protein MGC1223 (MGC1223), mRNA
NM_080920	Homo sapiens gamma-glutamyltransferase-like activity 4 (GGTLA4), mRNA
NM_021168	Homo sapiens RAR (RAS like GTPASE) like (RARL), mRNA
NM_080842	Homo sapiens hypothetical gene similar to gamma-glutamyltransferase-like
	activity 1 (LOC129026), mRNA
NM_031460	Homo sapiens potassium channel, subfamily K, member 17 (TASK-4)
	(KCNK17), mRNA
NM_033056	Homo sapiens protocadherin 15 (PCDH15), mRNA
NM_053283	Homo sapiens dermcidin (DCD), mRNA
NM_033518	Homo sapiens solute carrier family 38, member 5 (SLC38A5), mRNA
NM_021160	Homo sapiens HLA-B associated transcript 5 (BAT5), mRNA
NM 002279	Homo sapiens keratin, hair, acidic, 3B (KRTHA3B), mRNA
NM_004138	Homo sapiens keratin, hair, acidic, 3A (KRTHA3A), mRNA
NM 016310	Homo sapiens polymerase (RNA) III (DNA directed) polypeptide K (12.3 kD)
_	(POLR3K), mRNA
NM_031991	Homo sapiens polypyrimidine tract binding protein 1 (PTBP1), transcript variant
	3, mRNA
NM_031990	Homo sapiens polypyrimidine tract binding protein 1 (PTBP1), transcript variant
	2, mRNA
NM_002819	Homo sapiens polypyrimidine tract binding protein 1 (PTBP1), transcript variant
	1, mRNA
NM_030930	Homo sapiens unc-93 homolog B1 (C. elegans) (UNC93B1), mRNA
NM_022454	Homo sapiens SRY-related HMG-box transcription factor SOX17 (SOX17),
	mRNA
NM_004652	Homo sapiens ubiquitin specific protease 9, X chromosome (fat facets-like
	Drosophila) (USP9X), transcript variant 1, mRNA
NM_021906	Homo sapiens ubiquitin specific protease 9, X chromosome (fat facets-like
	Drosophila) (USP9X), transcript variant 2, mRNA
NM_022349	Homo sapiens membrane-spanning 4-domains, subfamily A, member 6A
	(MS4A6A), mRNA
NM_022122	Homo sapiens matrix metalloproteinase 27 (MMP27), mRNA
NM_006387	Homo sapiens calcium homeostasis endoplasmic reticulum protein (CHERP),
37.00.00	mRNA
NM_006918	Homo sapiens sterol-C5-desaturase (ERG3 delta-5-desaturase homolog, fungal)-
306 555 55	like (SC5DL), mRNA
NM_020151	Homo sapiens START domain containing 7 (STARD7), mRNA
NM_018976	Homo sapiens solute carrier family 38, member 2 (SLC38A2), mRNA
NM_013351	Homo sapiens T-box 21 (TBX21), mRNA
NM_006993	Homo sapiens nucleophosmin/nucleoplasmin, 3 (NPM3), mRNA
NM_002420	Homo sapiens transient receptor potential cation channel, subfamily M, member

1 (TRPMI), mRNA NM 007244 Homo sapiens proline rich 4 (lacrimal) (PROL4), mRNA NM 006758 Homo sapiens U2(RNU2) small nuclear RNA auxillary factor 1 (U2AF1), mRNA NM_006264 Homo sapiens protein tyrosine phosphatase, non-receptor type 13 (APO-1/CD95 (Fas)-associated phosphatase) (PTPN13), transcript variant 2, mRNA NM_006055 Homo sapiens LanC lantibiotic synthetase component C-like 1 (bacterial) (LANCL1), mRNA NM_00516 Homo sapiens regulator of G-protein signalling 19 interacting protein 1 (RGS19IP1), mRNA NM_005149 Homo sapiens T-box 19 (TBX19), mRNA NM_004231 Homo sapiens ATPase, vacuolar, 14 kD (ATP6S14), mRNA NM_000275 Homo sapiens atPrase, vacuolar, 14 kD (ATP6S14), mRNA NM_001384 Homo sapiens dipheria toxin resistance protein required for diphthamide biosynthesis-like 2 (S. cerevisiae) (DPH2L2), mRNA NM_000307 Homo sapiens serine (or cysteine) proteinase inhibitor, clade G (C1 inhibitor), member 1, (angiodema, hereditary) (SERPING1), mRNA NM_003307 Homo sapiens transient receptor potential cation channel, subfamily M, member 2 (TRPM2), mRNA NM_003807 Homo sapiens tumor necrosis factor (ligand) superfamily, member 14 (TNFSF14), mRNA NM_002984 Homo sapiens hemoglobin, theta 1 (HBQ1), mRNA NM_003931 Homo sapiens hemoglobin, theta 1 (HBQ1), mRNA NM_000531 Homo sapiens hemoglobin, alpha 1 (HBA1), mRNA NM_000517 Homo sapiens heparan sulfate 2-O-sulfotransferase 1 (HS2ST1), mRNA NM_001201 Homo sapiens heparan sulfate 2-O-sulfotransferase 1 (HS2ST1), mRNA NM_01213 Homo sapiens Phosphatidylcholine transfer protein (PCTP), mRNA NM_018960 Homo sapiens RNA binding protein (autoantigenic, hnRNA-associated with lethal yellow) (RALY), transcript variant 1, mRNA NM_016732 Homo sapiens Phosphatidylcholine transfer protein (RBMS3), mRNA NM_00184 Homo sapiens hemoglobin, gamma G (HBG2), mRNA NM_00184 Homo sapiens hemoglobin, gamma G (HBG2), mRNA NM_00184 Homo sapiens hemoglobin, gamma G (HBG2), mRNA NM_000589 Homo sapiens hemoglobin gamma G (HBG1), mRNA NM_000599 Homo sapiens hemoglobin gamma G (HBG2)		1 (mp.p) (1) p) [4
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CLANCLI), mRNA	_	(Fas)-associated phosphatase) (PTPN13), transcript variant 2, mRNA
CLANCL1), mRNA	NM 006055	Homo sapiens LanC lantibiotic synthetase component C-like 1 (bacterial)
RGS19IP1), mRNA Homo sapiens T-box 19 (TBX19), mRNA MM 004231 Homo sapiens ATPase, vacuolar, 14 kD (ATP6S14), mRNA MM 000275 Homo sapiens oculocutaneous albinism II (pink-eye dilution homolog, mouse) (OCA2), mRNA Homo sapiens diptheria toxin resistance protein required for diphthamide biosynthesis-like 2 (S. cerevisiae) (DPH2L2), mRNA MM 000062 Homo sapiens serine (or cysteine) proteinase inhibitor, clade G (C1 inhibitor), member 1, (angioedema, hereditary) (SERPING1), mRNA MM 003307 Homo sapiens transient receptor potential cation channel, subfamily M, member 2 (TRPM2), mRNA Homo sapiens tumor necrosis factor (ligand) superfamily, member 14 (TNFSF14), mRNA MM 002984 Homo sapiens small inducible cytokine A4 (SCYA4), mRNA MM 00105 Homo sapiens hemoglobin, alpha 1 (HBA1), mRNA MM 005331 Homo sapiens hemoglobin, alpha 1 (HBA1), mRNA MM 00558 Homo sapiens hemoglobin, alpha 1 (HBA1), mRNA MM 00558 Homo sapiens hemoglobin, alpha 2 (HBA2), mRNA MM 01262 Homo sapiens heparan sulfate 2-O-sulfotransferase 1 (HS2ST1), mRNA MM 018960 Homo sapiens phepatidylcholine transfer protein (PCTP), mRNA MM 018960 Homo sapiens phesididylcholine transfer protein (PCTP), mRNA MM 018960 Homo sapiens RNA binding protein (autoantigenic, hnRNP-associated with lethal yellow) (RALY), transcript variant 1, mRNA MM 01483 Homo sapiens RNA binding motif, single stranded interacting protein (RBMS3), mRNA MM 00184 Homo sapiens hemoglobin, gamma G (HBG2), mRNA MM 001330 Homo sapiens hemoglobin, paina G (HBG2), mRNA MM 005330 Homo sapiens hemoglobin, paina G (HBG2), mRNA MM 005331 Homo sapiens hemoglobin, paina G (HBG2), mRNA MM 005332 Homo sapiens RNA binding protein (autoantigenic, hnRNP-associated with lethal yellow) (RALY), transcript variant 2, mRNA MM 005331 Homo sapiens hemoglobin, paina G (HBG2), mRNA MM 005332 Homo sapiens hemoglobin, paina G (HBG2), mRNA MM 005334 Homo sapiens hemoglobin, paina G (HBG2), mRNA Homo sapiens hemoglobin, paina G ((LANCL1), mRNA
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NM 017807 Homo sapiens glycine N-methyltransferase (GNMT), mRNA NM 017807 Homo sapiens O-sialoglycoprotein endopeptidase (OSGEP), mRNA NM 016732 Homo sapiens RNA binding protein (autoantigenic, hnRNP-associated with lethal yellow) (RALY), transcript variant 1, mRNA NM 014483 Homo sapiens RNA binding motif, single stranded interacting protein (RBMS3), mRNA NM 012320 Homo sapiens lysophospholipase 3 (LYPLA3), mRNA NM 00184 Homo sapiens hemoglobin, gamma G (HBG2), mRNA NM 005330 Homo sapiens hemoglobin, epsilon 1 (HBE1), mRNA NM 007367 Homo sapiens RNA binding protein (autoantigenic, hnRNP-associated with lethal yellow) (RALY), transcript variant 2, mRNA NM 005332 Homo sapiens hemoglobin, zeta (HBZ), mRNA NM 005438 Homo sapiens FOS-like antigen 1 (FOSL1), mRNA NM 000158 Homo sapiens glucan (1,4-alpha-), branching enzyme 1 (glycogen branching enzyme, Andersen disease, glycogen storage disease type IV) (GBE1), mRNA	NM_012262	Homo sapiens heparan sulfate 2-O-sulfotransferase 1 (HS2ST1), mRNA
NM_017807 Homo sapiens O-sialoglycoprotein endopeptidase (OSGEP), mRNA NM_016732 Homo sapiens RNA binding protein (autoantigenic, hnRNP-associated with lethal yellow) (RALY), transcript variant 1, mRNA NM_014483 Homo sapiens RNA binding motif, single stranded interacting protein (RBMS3), mRNA NM_012320 Homo sapiens lysophospholipase 3 (LYPLA3), mRNA NM_000184 Homo sapiens hemoglobin, gamma G (HBG2), mRNA NM_005330 Homo sapiens hemoglobin, epsilon 1 (HBE1), mRNA NM_007367 Homo sapiens RNA binding protein (autoantigenic, hnRNP-associated with lethal yellow) (RALY), transcript variant 2, mRNA NM_005332 Homo sapiens hemoglobin, zeta (HBZ), mRNA NM_005438 Homo sapiens FOS-like antigen 1 (FOSL1), mRNA NM_000158 Homo sapiens glucan (1,4-alpha-), branching enzyme 1 (glycogen branching enzyme, Andersen disease, glycogen storage disease type IV) (GBE1), mRNA	NM_021213	Homo sapiens phosphatidylcholine transfer protein (PCTP), mRNA
NM_016732 Homo sapiens RNA binding protein (autoantigenic, hnRNP-associated with lethal yellow) (RALY), transcript variant 1, mRNA NM_014483 Homo sapiens RNA binding motif, single stranded interacting protein (RBMS3), mRNA NM_012320 Homo sapiens lysophospholipase 3 (LYPLA3), mRNA NM_000184 Homo sapiens hemoglobin, gamma G (HBG2), mRNA NM_005330 Homo sapiens hemoglobin, epsilon 1 (HBE1), mRNA NM_007367 Homo sapiens RNA binding protein (autoantigenic, hnRNP-associated with lethal yellow) (RALY), transcript variant 2, mRNA NM_005332 Homo sapiens hemoglobin, zeta (HBZ), mRNA NM_005438 Homo sapiens FOS-like antigen 1 (FOSL1), mRNA NM_000158 Homo sapiens glucan (1,4-alpha-), branching enzyme 1 (glycogen branching enzyme, Andersen disease, glycogen storage disease type IV) (GBE1), mRNA	NM_018960	Homo sapiens glycine N-methyltransferase (GNMT), mRNA
lethal yellow) (RALY), transcript variant 1, mRNA NM_014483 Homo sapiens RNA binding motif, single stranded interacting protein (RBMS3), mRNA NM_012320 Homo sapiens lysophospholipase 3 (LYPLA3), mRNA NM_000184 Homo sapiens hemoglobin, gamma G (HBG2), mRNA NM_005330 Homo sapiens hemoglobin, epsilon 1 (HBE1), mRNA NM_007367 Homo sapiens RNA binding protein (autoantigenic, hnRNP-associated with lethal yellow) (RALY), transcript variant 2, mRNA NM_005332 Homo sapiens hemoglobin, zeta (HBZ), mRNA NM_005438 Homo sapiens FOS-like antigen 1 (FOSL1), mRNA NM_000158 Homo sapiens glucan (1,4-alpha-), branching enzyme 1 (glycogen branching enzyme, Andersen disease, glycogen storage disease type IV) (GBE1), mRNA	NM_017807	Homo sapiens O-sialoglycoprotein endopeptidase (OSGEP), mRNA
NM_014483 Homo sapiens RNA binding motif, single stranded interacting protein (RBMS3), mRNA NM_012320 Homo sapiens lysophospholipase 3 (LYPLA3), mRNA NM_000184 Homo sapiens hemoglobin, gamma G (HBG2), mRNA NM_005330 Homo sapiens hemoglobin, epsilon 1 (HBE1), mRNA NM_007367 Homo sapiens RNA binding protein (autoantigenic, hnRNP-associated with lethal yellow) (RALY), transcript variant 2, mRNA NM_005332 Homo sapiens hemoglobin, zeta (HBZ), mRNA NM_005438 Homo sapiens FOS-like antigen 1 (FOSL1), mRNA NM_000158 Homo sapiens glucan (1,4-alpha-), branching enzyme 1 (glycogen branching enzyme, Andersen disease, glycogen storage disease type IV) (GBE1), mRNA	NM_016732	
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NM_007367 Homo sapiens RNA binding protein (autoantigenic, hnRNP-associated with lethal yellow) (RALY), transcript variant 2, mRNA NM_005332 Homo sapiens hemoglobin, zeta (HBZ), mRNA NM_005438 Homo sapiens FOS-like antigen 1 (FOSL1), mRNA NM_000158 Homo sapiens glucan (1,4-alpha-), branching enzyme 1 (glycogen branching enzyme, Andersen disease, glycogen storage disease type IV) (GBE1), mRNA	NM_000184	Homo sapiens hemoglobin, gamma G (HBG2), mRNA
lethal yellow) (RALY), transcript variant 2, mRNA NM_005332 Homo sapiens hemoglobin, zeta (HBZ), mRNA NM_005438 Homo sapiens FOS-like antigen 1 (FOSL1), mRNA NM_000158 Homo sapiens glucan (1,4-alpha-), branching enzyme 1 (glycogen branching enzyme, Andersen disease, glycogen storage disease type IV) (GBE1), mRNA	NM_005330	Homo sapiens hemoglobin, epsilon 1 (HBE1), mRNA
lethal yellow) (RALY), transcript variant 2, mRNA NM_005332 Homo sapiens hemoglobin, zeta (HBZ), mRNA NM_005438 Homo sapiens FOS-like antigen 1 (FOSL1), mRNA NM_000158 Homo sapiens glucan (1,4-alpha-), branching enzyme 1 (glycogen branching enzyme, Andersen disease, glycogen storage disease type IV) (GBE1), mRNA	NM_007367	Homo sapiens RNA binding protein (autoantigenic, hnRNP-associated with
NM_005438 Homo sapiens FOS-like antigen 1 (FOSL1), mRNA NM_000158 Homo sapiens glucan (1,4-alpha-), branching enzyme 1 (glycogen branching enzyme, Andersen disease, glycogen storage disease type IV) (GBE1), mRNA		
NM_000158 Homo sapiens glucan (1,4-alpha-), branching enzyme 1 (glycogen branching enzyme, Andersen disease, glycogen storage disease type IV) (GBE1), mRNA	NM_005332	Homo sapiens hemoglobin, zeta (HBZ), mRNA
NM_000158 Homo sapiens glucan (1,4-alpha-), branching enzyme 1 (glycogen branching enzyme, Andersen disease, glycogen storage disease type IV) (GBE1), mRNA	NM_005438	Homo sapiens FOS-like antigen 1 (FOSL1), mRNA
enzyme, Andersen disease, glycogen storage disease type IV) (GBE1), mRNA	NM_000158	Homo sapiens glucan (1,4-alpha-), branching enzyme 1 (glycogen branching
NM 000559 Homo saniens hemoglobin, gamma A (HBG1), mRNA		enzyme, Andersen disease, glycogen storage disease type IV) (GBE1), mRNA
	NM_000559	Homo sapiens hemoglobin, gamma A (HBG1), mRNA
NG_000007 Homo sapiens genomic beta globin region (HBB@) on chromosome 11	NG_000007	
NG_000006 Homo sapiens genomic alpha globin region (HBA@) on chromosome 16		Homo sapiens genomic alpha globin region (HBA@) on chromosome 16
NM_030964 Homo sapiens sprouty homolog 4 (Drosophila) (SPRY4), mRNA		
NM_021181 Homo sapiens 19A24 protein (CRACC), mRNA		Homo sapiens 19A24 protein (CRACC), mRNA
NM_004654 Homo sapiens ubiquitin specific protease 9, Y chromosome (fat facets-like Drosophila) (USP9Y), mRNA	NM_004654	Homo sapiens ubiquitin specific protease 9, Y chromosome (fat facets-like
NM_018518 Homo sapiens MCM10 minichromosome maintenance deficient 10 (S.	NM 018518	

	T \ 2.50(16) D111
	cerevisiae) (MCM10), mRNA
NM_018593	Homo sapiens solute carrier family 16 (monocarboxylic acid transporters),
	member 10 (SLC16A10), mRNA
NM_018240	Homo sapiens kin of IRRE like (Drosophila) (KIRREL), mRNA
NM_016004	Homo sapiens chromosome 20 open reading frame 9 (C20orf9), mRNA
NM_006841	Homo sapiens solute carrier family 38, member 3 (SLC38A3), mRNA
NM_003725	Homo sapiens oxidative 3 alpha hydroxysteroid dehydrogenase; retinol
	dehydrogenase; 3-hydroxysteroid epimerase (RODH), mRNA
NG_000009	Homo sapiens genomic small histone family cluster (HFS@) on chromosome 6
NM_080878	Homo sapiens endothelial lectin HL-2 (HL-2), mRNA
NM_080876	Homo sapiens protein phosphatase (SKRP1), mRNA
NM_080874	Homo sapiens ankyrin repeat and SOCS box-containing 5 (ASB5), mRNA
NM_080873	Homo sapiens ankyrin repeat and SOCS box-containing 11 (ASB11), mRNA
NM_080872	Homo sapiens KIAA1777 protein (Unc5h4), mRNA
NM_080867	Homo sapiens suppressor of cytokine signalling 4 (SOCS4), mRNA
NM_080864	Homo sapiens relaxin 3 (H3) (RLN3), mRNA
NM_080863	Homo sapiens ankyrin repeat and SOCS box-containing 16 (ASB16), mRNA
NM_080862	Homo sapiens SPRY domain-containing SOCS box protein SSB-4 (SSB-4), mRNA
NM_080861	Homo sapiens SPRY domain-containing SOCS box protein SSB-3 (SSB-3), mRNA
NM 080860	Homo sapiens testes specific A2 homolog (mouse) (TSGA2), mRNA
NM 016150	Homo sapiens ankyrin repeat and SOCS box-containing 2 (ASB2), mRNA
NM 016127	Homo sapiens hypothetical protein MGC8721 (MGC8721), mRNA
NM 004170	Homo sapiens solute carrier family 1 (neuronal/epithelial high affinity glutamate
11112_00 1170	transporter, system Xag), member 1 (SLC1A1), nuclear gene encoding
	mitochondrial protein, mRNA
NM 017611	Homo sapiens hypothetical protein DKFZp762A227 (DKFZp762A227), mRNA
NM_025220	Homo sapiens a disintegrin and metalloproteinase domain 33 (ADAM33), mRNA
NM_018548	Homo sapiens down-regulated in lung cancer (HLCDGP1), mRNA
NM_080740	Homo sapiens similar to Ovis aries Y chromosome repeat region OY11.1 (3'OY11.1), mRNA
NM 012163	Homo sapiens F-box and leucine-rich repeat protein 9 (FBXL9), mRNA
NM 012304	Homo sapiens F-box and leucine-rich repeat protein 7 (FBXL7), mRNA
NM 012160	Homo sapiens F-box and leucine-rich repeat protein 4 (FBXL4), mRNA
NM 012159	Homo sapiens F-box and leucine-rich repeat protein 3B (FBXL3B), mRNA
NM 012158	Homo sapiens F-box and leucine-rich repeat protein 3A (FBXL3A), mRNA
NM 012157	Homo sapiens F-box and leucine-rich repeat protein 2 (FBXL2), mRNA
NM_024555	Homo sapiens F-box and leucine-rich repeat protein 2 (FBXL2), mkNA Homo sapiens F-box and leucine-rich repeat protein 6 (FBXL6), transcript
14MI_024333	variant 2, mRNA
NM 012162	Homo sapiens F-box and leucine-rich repeat protein 6 (FBXL6), transcript
NW1_012102	variant 1, mRNA
NM 033535	Homo sapiens F-box and leucine-rich repeat protein 5 (FBXL5), transcript
IAIM_022222	
NDA 012161	variant 2, mRNA Homo sapiens F-box and leucine-rich repeat protein 5 (FBXL5), transcript
NM_012161	variant 1, mRNA
NIM 002279	
NM 002278	Homo sapiens keratin, hair, acidic, 2 (KRTHA2), mRNA
NM_033285	Homo sapiens tumor protein p53 inducible nuclear protein 1 (TP53INP1), mRNA
NM_002277	Homo sapiens keratin, hair, acidic, 1 (KRTHA1), mRNA
NM 032994	Homo sapiens Williams Beuren syndrome chromosome region 14 (WBSCR14),

	L
127.6.000.64	transcript variant 5, mRNA
NM_032954	Homo sapiens Williams Beuren syndrome chromosome region 14 (WBSCR14),
3 7 7 7 7 7 7 7 7 7 7 7 7 7 7 7 7 7 7 7	transcript variant 4, mRNA
NM_032953	Homo sapiens Williams Beuren syndrome chromosome region 14 (WBSCR14),
	transcript variant 3, mRNA
NM_032952	Homo sapiens Williams Beuren syndrome chromosome region 14 (WBSCR14),
	transcript variant 2, mRNA
NM_032951	Homo sapiens Williams Beuren syndrome chromosome region 14 (WBSCR14),
	transcript variant 1, mRNA
NG_000008	Homo sapiens genomic cytochrome P450, subfamily IIA (phenobarbital-
	inducible) (CYP2A) on chromosome 19
NM_030809	Homo sapiens chromosome 12 open reading frame 22 (C12orf22), mRNA
NM_004426	Homo sapiens early development regulator 1 (polyhomeotic 1 homolog) (EDR1),
	mRNA
NM_020244	Homo sapiens choline phosphotransferase 1 (CHPT1), mRNA
NM_019074	Homo sapiens delta-like 4 (Drosophila) (DLL4), mRNA
NM_018990	Homo sapiens chromosome X open reading frame 9 (CXorf9), mRNA
NM 017833	Homo sapiens chromosome 21 open reading frame 55 (C21orf55), mRNA
NM 018255	Homo sapiens elongator protein 2 (ELP2), mRNA
NM 014096	Homo sapiens hypothetical protein DKFZp762A227 (DKFZp762A227), mRNA
NM 014927	Homo sapiens connector enhancer of KSR2 (CNK2), mRNA
NM 012164	Homo sapiens F-box and WD-40 domain protein 2 (FBXW2), mRNA
NM 012247	Homo sapiens selenium donor protein (SPS), mRNA
NM 012165	Homo sapiens F-box and WD-40 domain protein 3 (FBXW3), mRNA
NM 007198	Homo sapiens proline synthetase co-transcribed homolog (bacterial) (PROSC),
14141_007130	mRNA
NM_006011	Homo sapiens sialyltransferase 8B (alpha-2, 8-sialytransferase) (SIAT8B),
-	mRNA
NM 005674	Homo sapiens zinc finger protein 239 (ZNF239), mRNA
NM 001364	Homo sapiens discs, large homolog 2, chapsyn-110 (Drosophila) (DLG2),
-	mRNA
NM 000646	Homo sapiens amylo-1, 6-glucosidase, 4-alpha-glucanotransferase (glycogen
_	debranching enzyme, glycogen storage disease type III) (AGL), transcript variant
	6, mRNA
NM 000645	Homo sapiens amylo-1, 6-glucosidase, 4-alpha-glucanotransferase (glycogen
-	debranching enzyme, glycogen storage disease type III) (AGL), transcript variant
	5, mRNA
NM 000644	Homo sapiens amylo-1, 6-glucosidase, 4-alpha-glucanotransferase (glycogen
_	debranching enzyme, glycogen storage disease type III) (AGL), transcript variant
	2, mRNA
NM 000643	Homo sapiens amylo-1, 6-glucosidase, 4-alpha-glucanotransferase (glycogen
_	debranching enzyme, glycogen storage disease type III) (AGL), transcript variant
	3, mRNA
NM 000642	Homo sapiens amylo-1, 6-glucosidase, 4-alpha-glucanotransferase (glycogen
	debranching enzyme, glycogen storage disease type III) (AGL), transcript variant
	1, mRNA
NM 000028	Homo sapiens amylo-1, 6-glucosidase, 4-alpha-glucanotransferase (glycogen
	debranching enzyme, glycogen storage disease type III) (AGL), transcript variant
	4, mRNA
NM 080831	Homo sapiens chromosome 20 open reading frame 87 (C20orf87), mRNA
NM_080825	Homo sapiens chromosome 20 open reading frame 144 (C20orf144), mRNA
NM 080823	Homo sapiens chromosome 20 open reading frame 148 (C20orf148), mRNA
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NM_017662	Homo sapiens transient receptor potential cation channel, subfamily M, member 6 (TRPM6), mRNA
NM_080744	Homo sapiens scavenger receptor cysteine rich domain containing, group B (4 domains) (SRCRB4D), mRNA
NM_000493	Homo sapiens collagen, type X, alpha 1(Schmid metaphyseal chondrodysplasia) (COL10A1), mRNA
NM_057096	Homo sapiens cytochrome P450 polypeptide 43 (CYP3A43), transcript variant 3, mRNA
NM 014578	Homo sapiens ras homolog gene family, member D (ARHD), mRNA
NM_020708	Homo sapiens solute carrier family 12, (potassium-chloride transporter) member 5 (SLC12A5), mRNA
NM 016093	Homo sapiens ribosomal protein L26-like 1 (RPL26L1), mRNA
NM_057095	Homo sapiens cytochrome P450 polypeptide 43 (CYP3A43), transcript variant 2, mRNA
NM_022820	Homo sapiens cytochrome P450 polypeptide 43 (CYP3A43), transcript variant 1, mRNA
NM_052969	Homo sapiens ribosomal protein L39-like (RPL39L), mRNA
NM_052970	Homo sapiens chromosome 20 open reading frame 60 (C20orf60), mRNA
NM_052865	Homo sapiens chromosome 20 open reading frame 72 (C20orf72), mRNA
NM_021029	Homo sapiens ribosomal protein L36a (RPL36A), mRNA
NM_001001	Homo sapiens ribosomal protein L36a-like (RPL36AL), mRNA
NM_033645	Homo sapiens F-box and WD-40 domain protein 1B (FBXW1B), transcript variant 1, mRNA
NM_033644	Homo sapiens F-box and WD-40 domain protein 1B (FBXW1B), transcript variant 2, mRNA
NM_012300	Homo sapiens F-box and WD-40 domain protein 1B (FBXW1B), transcript variant 3, mRNA
NM 022760	Homo sapiens chromosome 20 open reading frame 81 (C20orf81), mRNA
NM_014958	Homo sapiens Rho guanine nucleotide exchange factor (GEF) 15 (ARHGEF15), mRNA
NM 021810	Homo sapiens cadherin-like 26 (CDH26), mRNA
NM_030876	Homo sapiens olfactory receptor, family 5, subfamily V, member 1 (OR5V1), mRNA
NM_031232	Homo sapiens amyloid beta (A4) precursor protein-binding, family A, member 2 binding protein (APBA2BP), transcript variant 2, mRNA
NM_031231	Homo sapiens amyloid beta (A4) precursor protein-binding, family A, member 2 binding protein (APBA2BP), transcript variant 1, mRNA
NM_032554	Homo sapiens G protein-coupled receptor 81 (GPR81), mRNA
NM_006462	Homo sapiens chromosome 20 open reading frame 18 (C20orf18), transcript variant 1, mRNA
NM_031229	Homo sapiens chromosome 20 open reading frame 18 (C20orf18), transcript variant 2, mRNA
NM_031228	Homo sapiens chromosome 20 open reading frame 18 (C20orf18), transcript variant 3, mRNA
NM_031227	Homo sapiens chromosome 20 open reading frame 18 (C20orf18), transcript variant 4, mRNA
NM_031424	Homo sapiens chromosome 20 open reading frame 55 (C20orf55), mRNA
NM_000518	Homo sapiens hemoglobin, beta (HBB), mRNA
NM_030959	Homo sapiens olfactory receptor, family 12, subfamily D, member 3 (OR12D3), mRNA
NM_018661	Homo sapiens defensin, beta 3 (DEFB3), mRNA
NM_022487	Homo sapiens DNA cross-link repair 1C (PSO2 homolog, S. cerevisiae)

	(DOLDETO) DNA
3D 6 000000	(DCLRE1C), mRNA Homo sapiens chromosome 20 open reading frame 51 (C20orf51), mRNA
NM_022099	Homo sapiens alcohol dehydrogenase IB (class I), beta polypeptide (ADH1B),
NM_000668	mRNA
NM 021943	Homo sapiens testis expressed sequence 27 (TEX27), mRNA
NM 021640	Homo sapiens chromosome 12 open reading frame 10 (C12orf10), mRNA
NM 021215	Homo sapiens chromosome 20 open reading frame 77 (C20orf77), mRNA
NM_012141	Homo sapiens DEAD/H (Asp-Glu-Ala-Asp/His) box polypeptide 26 (DDX26), mRNA
NM 021225	Homo sapiens proline-rich 1 (PROL1), mRNA
NM 006508	Homo sapiens regenerating islet-derived-like, pancreatic stone protein-like,
_	pancreatic thread protein-like (rat) (REGL), mRNA
NM_020356	Homo sapiens chromosome 20 open reading frame 32 (C20orf32), mRNA
NM 020369	Homo sapiens fascin homolog 3, actin-bundling protein, testicular
_	(Strongylocentrotus purpuratus) (FSCN3), mRNA
NM 020145	Homo sapiens SH3-domain GRB2-like endophilin B2 (SH3GLB2), mRNA
NM_020125	Homo sapiens BCM-like membrane protein precursor (BLAME), mRNA
NM_019025	Homo sapiens chromosome 20 open reading frame 16 (C20orf16), mRNA
NM_018679	Homo sapiens t-complex 11 (mouse) (TCP11), mRNA
NM_017589	Homo sapiens B-cell translocation gene 4 (BTG4), mRNA
NM_018692	Homo sapiens chromosome 20 open reading frame 17 (C20orf17), mRNA
NM_018697	Homo sapiens LanC lantibiotic synthetase component C-like 2 (bacterial) (LANCL2), mRNA
NM_018677	Homo sapiens acetyl-Coenzyme A synthetase 2 (ADP forming) (ACAS2), mRNA
NM 018431	Homo sapiens chromosome 20 open reading frame 180 (C20orf180), mRNA
NM 018725	Homo sapiens interleukin 17B receptor (IL17BR), mRNA
NM 018474	Homo sapiens chromosome 20 open reading frame 19 (C20orf19), mRNA
NM 018478	Homo sapiens chromosome 20 open reading frame 35 (C20orf35), mRNA
NM 017896	Homo sapiens chromosome 20 open reading frame 11 (C20orf11), mRNA
NM 017874	Homo sapiens chromosome 20 open reading frame 27 (C20orf27), mRNA
NM 017859	Homo sapiens uridine kinase-like 1 (URKL1), mRNA
NM_017798	Homo sapiens chromosome 20 open reading frame 21 (C20orf21), mRNA
NM_017789	Homo sapiens sema domain, immunoglobulin domain (Ig), transmembrane domain (TM) and short cytoplasmic domain, (semaphorin) 4C (SEMA4C), mRNA
NM_017714	Homo sapiens chromosome 20 open reading frame 13 (C20orf13), mRNA
NM_017671	Homo sapiens chromosome 20 open reading frame 42 (C20orf42), mRNA
NM_018384	Homo sapiens immune associated nucleotide 4 like 1 (mouse) (IAN4L1), mRNA
NM_018354	Homo sapiens chromosome 20 open reading frame 46 (C20orf46), mRNA
NM_018347	Homo sapiens chromosome 20 open reading frame 29 (C20orf29), mRNA
NM_018327	Homo sapiens chromosome 20 open reading frame 38 (C20orf38), mRNA
NM_018282	Homo sapiens paraspeckle protein 1 (PSP1), mRNA
NM_018270	Homo sapiens chromosome 20 open reading frame 20 (C20orf20), mRNA
NM_018257	Homo sapiens chromosome 20 open reading frame 36 (C20orf36), mRNA
NM_018197	Homo sapiens zinc finger protein 64 homolog (mouse) (ZFP64), mRNA
NM_018010	Homo sapiens estrogen-related receptor beta like 1 (ESRRBL1), mRNA
NM_017446	Homo sapiens mitochondrial ribosomal protein L39 (MRPL39), mRNA
NM_017429	Homo sapiens beta-carotene 15, 15'-dioxygenase (BCDO), mRNA
NM_016082	Homo sapiens chromosome 20 open reading frame 34 (C20orf34), mRNA
NM_016610	Homo sapiens toll-like receptor 8 (TLR8), mRNA
NM_016009	Homo sapiens SH3-domain GRB2-like endophilin B1 (SH3GLB1), mRNA

NM_016408	Homo sapiens chromosome 20 open reading frame 34 (C20orf34), mRNA
NM_016407	Homo sapiens chromosome 20 open reading frame 43 (C20orf43), mRNA
NM_016319	Homo sapiens COP9 constitutive photomorphogenic homolog subunit 7A (Arabidopsis) (COPS7A), mRNA
NM 015985	Homo sapiens angiopoietin 4 (ANGPT4), mRNA
NM 015834	Homo sapiens adenosine deaminase, RNA-specific, B1 (RED1 homolog rat)
-	(ADARB1), transcript variant DRADA2c, mRNA
NM_015833	Homo sapiens adenosine deaminase, RNA-specific, B1 (RED1 homolog rat)
	(ADARB1), transcript variant DRABA2b, mRNA
NM_014036	Homo sapiens BCM-like membrane protein precursor (BLAME), mRNA
NM_014012	Homo sapiens RAS (RAD and GEM)-like GTP-binding (REM), mRNA
NM_014841	Homo sapiens synaptosomal-associated protein, 91 kD homolog (mouse) (SNAP91), mRNA
NM_014795	Homo sapiens zinc finger homeobox 1b (ZFHX1B), mRNA
NM 015313	Homo sapiens Rho guanine nucleotide exchange factor (GEF) 12 (ARHGEF12),
	mRNA
NM_014784	Homo sapiens Rho guanine nucleotide exchange factor (GEF) 11 (ARHGEF11), mRNA
NM_014862	Homo sapiens aryl-hydrocarbon receptor nuclear translocator 2 (ARNT2),
1111_014802	mRNA
NM_014054	Homo sapiens chromosome 20 open reading frame 40 (C20orf40), mRNA
NM_015629	Homo sapiens PRP31 pre-mRNA processing factor 31 homolog (yeast)
	(PRPF31), mRNA
NM_015417	Homo sapiens chromosome 20 open reading frame 28 (C20orf28), mRNA
NM_014625	Homo sapiens nephrosis 2, idiopathic, steroid-resistant (podocin) (NPHS2), mRNA
NM 014592	Homo sapiens Kv channel interacting protein 1 (KCNIP1), mRNA
NM_014140	Homo sapiens SWI/SNF related, matrix associated, actin dependent regulator of chromatin, subfamily a-like 1 (SMARCAL1), mRNA
NM 013442	Homo sapiens stomatin (EPB72)-like 2 (STOML2), mRNA
NM 013248	Homo sapiens NUTF-like export factor 1 (NXT1), mRNA
NM_013316	Homo sapiens CCR4-NOT transcription complex, subunit (CNOT4), mRNA
NM 013348	Homo sapiens potassium inwardly-rectifying channel, subfamily J, member 14
-	(KCNJ14), mRNA
NM_013279	Homo sapiens chromosome 11 open reading frame 9 (C11orf9), mRNA
NM_012418	Homo sapiens fascin homolog 2, actin-bundling protein, retinal
	(Strongylocentrotus purpuratus) (FSCN2), mRNA
NM_012201	Homo sapiens golgi apparatus protein 1 (GLG1), mRNA
NM_000519	Homo sapiens hemoglobin, delta (HBD), mRNA
NM_006999	Homo sapiens polymerase (DNA directed) sigma (POLS), mRNA
NM_006719	Homo sapiens actin binding LIM protein (ABLIM), transcript variant ABLIM-m, mRNA
NM_002313	Homo sapiens actin binding LIM protein (ABLIM), transcript variant ABLIM-1, mRNA
NM 007238	
NM 007184	Homo sapiens peroxisomal membrane protein 4 (24kD) (PXMP4), mRNA Homo sapiens nischarin (NISCH), mRNA
NM_006720	Homo sapiens actin binding LIM protein (ABLIM), transcript variant ABLIM-s,
	mRNA
NM_007026	Homo sapiens dual specificity phosphatase 14 (DUSP14), mRNA
NM_006837	Homo sapiens COP9 constitutive photomorphogenic homolog subunit 5 (Arabidopsis) (COPS5), mRNA
NM_006614	Homo sapiens cell adhesion molecule with homology to L1CAM (close homolog
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	CT 1) (CTT 1) -DNA
ND (00 (410	of L1) (CHL1), mRNA Homo sapiens HIV-1 Tat interactive protein 2, 30 kD (HTATIP2), mRNA
NM_006410	Homo sapiens HIV-1 1at interactive protein 2, 50 kD (HIATIF2), mkiva
NM_006432	Homo sapiens Niemann-Pick disease, type C2 (NPC2), mRNA
NM_006348	Homo sapiens golgi transport complex 1 (90 kD subunit) (GOLTC1), mRNA
NM_006408	Homo sapiens anterior gradient 2 homolog (Xenepus laevis) (AGR2), mRNA
NM_006106	Homo sapiens Yes-associated protein 1, 65 kD (YAP1), mRNA
NM_006096	Homo sapiens N-myc downstream regulated gene 1 (NDRG1), mRNA
NM_006071	Homo sapiens polycystic kidney disease (polycystin) and REJ (sperm receptor for egg jelly homolog, sea urchin)-like (PKDREJ), mRNA
AD (000000	Homo sapiens caspase recruitment domain family, member 4 (CARD4), mRNA
NM_006092	
NM_005748	Homo sapiens YY1 associated factor 2 (YAF2), mRNA Homo sapiens uronyl-2-sulfotransferase (UST), mRNA
NM_005715	Homo sapiens SA hypertension-associated homolog (rat) (SAH), mRNA
NM_005622	Homo sapiens SA hyperension-associated homolog (lat) (SAH), higher
NM_005733	Homo sapiens RAB6 interacting, kinesin-like (rabkinesin6) (RAB6KIFL), mRNA
NM_005668	Homo sapiens sialyltransferase 8D (alpha-2, 8-polysialytransferase) (SIAT8D), mRNA
NM_005606	Homo sapiens legumain (LGMN), mRNA
NM_004649	Homo sapiens chromosome 21 open reading frame 33 (C21orf33), mRNA
NM_005469	Homo sapiens peroxisomal acyl-CoA thioesterase (PTE1), mRNA
NM_005180	Homo sapiens B lymphoma Mo-MLV insertion region (mouse) (BMI1), mRNA
NM 005108	Homo sapiens xylulokinase homolog (H. influenzae) (XYLB), mRNA
NM 004610	Homo sapiens t-complex 10 (mouse) (TCP10), mRNA
NM_004579	Homo sapiens mitogen-activated protein kinase kinase kinase kinase 2 (MAP4K2), mRNA
NM_004086	Homo sapiens coagulation factor C homolog, cochlin (Limulus polyphemus) (COCH), mRNA
NM 004273	Homo sapiens carbohydrate (chondroitin 6) sulfotransferase 3 (CHST3), mRNA
NM_004902	Homo sapiens RNA-binding region (RNP1, RRM) containing 2 (RNPC2), mRNA
NM_004353	Homo sapiens serine (or cysteine) proteinase inhibitor, clade H (heat shock
27.6 004017	protein 47), member 1, (collagen binding protein 1) (SERPINH1), mRNA
NM_004317	Homo sapiens arsA arsenite transporter, ATP-binding, homolog 1 (bacterial) (ASNA1), mRNA
NM_001247	Homo sapiens ectonucleoside triphosphate diphosphohydrolase 6 (putative function) (ENTPD6), mRNA
NM_003831	Homo sapiens sudD suppressor of bimD6 homolog (A. nidulans) (SUDD), mRNA
NM 003143	Homo sapiens single-stranded DNA binding protein (SSBP1), mRNA
NM_003098	Homo sapiens syntrophin, alpha 1 (dystrophin-associated protein A1, 59kD, acidic component) (SNTA1), mRNA
NM_003034	Homo sapiens sialyltransferase 8A (alpha-N-acetylneuraminate/alpha-2,8-
NR 6 600000	sialytransferase, GD3 synthase) (SIAT8A), mRNA
NM_003028	Homo sapiens SHB (Src homology 2 domain-containing) adaptor protein B (SHB), mRNA
NM_003579	Homo sapiens RAD54-like (S. cerevisiae) (RAD54L), mRNA
NM_002669	Homo sapiens pleiotropic regulator 1 (PRL1homolog, Arabidopsis) (PLRG1), mRNA
NM_000139	Homo sapiens membrane-spanning 4-domains, subfamily A, member 1 (MS4A2), mRNA
NM 003836	Homo sapiens delta-like 1 homolog (Drosophila) (DLK1), mRNA
NM 003653	Homo sapiens COP9 constitutive photomorphogenic homolog subunit 3

	(Arabidopsis) (COPS3), mRNA
NM 000083	Homo sapiens chloride channel 1, skeletal muscle (Thomsen disease, autosomal
1414_000003	dominant) (CLCN1), mRNA
NM_000691	Homo sapiens aldehyde dehydrogenase 3 family, memberA1 (ALDH3A1),
1442_000051	mRNA
NM 001112	Homo sapiens adenosine deaminase, RNA-specific, B1 (RED1 homolog rat)
	(ADARB1), transcript variant DRADA2a, mRNA
NM 004370	Homo sapiens collagen, type XII, alpha 1 (COL12A1), transcript variant long,
_	mRNA
NM 080645	Homo sapiens collagen, type XII, alpha 1 (COL12A1), transcript variant short,
	mRNA
NM_080681	Homo sapiens collagen, type XI, alpha 2 (COL11A2), transcript variant 2,
	mRNA
NM_080680	Homo sapiens collagen, type XI, alpha 2 (COL11A2), transcript variant 1,
	mRNA
NM_080679	Homo sapiens collagen, type XI, alpha 2 (COL11A2), transcript variant 3,
	mRNA
NM_003593	Homo sapiens winged-helix nude (WHN), mRNA
NM_000638	Homo sapiens vitronectin (serum spreading factor, somatomedin B, complement
377 C 000 C00	S-protein) (VTN), mRNA
NM_080682	Homo sapiens vascular cell adhesion molecule 1 (VCAM1), transcript variant 2,
NM_001078	mRNA Homo sapiens vascular cell adhesion molecule 1 (VCAM1), transcript variant 1,
141A1_0010\2	mRNA
NM_006115	Homo sapiens preferentially expressed antigen in melanoma (PRAME), mRNA
NM 000175	Homo sapiens glucose phosphate isomerase (GPI), mRNA
NM 020526	Homo sapiens EphA8 (EPHA8), mRNA
NM 002109	Homo sapiens histidyl-tRNA synthetase (HARS), mRNA
NM 012208	Homo sapiens histidyl-tRNA synthetase-like (HARSL), mRNA
NM 004608	Homo sapiens T-box 6 (TBX6), transcript variant 1, mRNA
NM_080758	Homo sapiens T-box 6 (TBX6), transcript variant 2, mRNA
NM_080718	Homo sapiens T-box 5 (TBX5), transcript variant 2, mRNA
NM 080717	Homo sapiens T-box 5 (TBX5), transcript variant 3, mRNA
NM 000192	Homo sapiens T-box 5 (TBX5), transcript variant 1, mRNA
NM 080832	Homo sapiens poly(A) binding protein, cytoplasmic 5 (PABPC5), mRNA
NM 080824	Homo sapiens chromosome 20 open reading frame 106 (C20orf106), mRNA
NM 080822	Homo sapiens candidate tumor suppressor OVCA2 (OVCA2), mRNA
NM 080821	Homo sapiens chromosome 20 open reading frame 108 (C20orf108), mRNA
NM 080820	Homo sapiens chromosome 20 open reading frame 88 (C20orf88), mRNA
NM_080818	Homo sapiens G protein-coupled receptor 80 (GPR80), mRNA
NM_080817	Homo sapiens G protein-coupled receptor 82 (GPR82), mRNA
NM_080794	Homo sapiens mitochondrial ribosomal protein L39 (MRPL39), mRNA
NM_020973	Homo sapiens cytosolic beta-glucosidase (GLUC), mRNA
NM_054112	Homo sapiens chromosome 20 open reading frame 63 (C20orf63), mRNA
NM_052951	Homo sapiens chromosome 20 open reading frame 167 (C20orf167), mRNA
NM_014145	Homo sapiens chromosome 20 open reading frame 30 (C20orf30), mRNA
NM_033409	Homo sapiens chromosome 20 open reading frame 54 (C20orf54), mRNA
NM_032013	Homo sapiens NDRG family member 3 (NDRG3), mRNA
NM_032109	Homo sapiens orthopedia homolog (Drosophila) (OTP), mRNA
NM_024021	Homo sapiens membrane-spanning 4-domains, subfamily A, member 4
	(MS4A4A), mRNA
NM_022910	Homo sapiens NDRG family member 4 (NDRG4), mRNA

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NM_025206	Homo sapiens fer-1-like 4 (C. elegans) (FER1L4), mRNA
NM_024960	Homo sapiens chromosome 20 open reading frame 48 (C20orf48), mRNA
NM_024893	Homo sapiens chromosome 20 open reading frame 39 (C20orf39), mRNA
NM_024299	Homo sapiens chromosome 20 open reading frame 149 (C20orf149), mRNA
NM_024077	Homo sapiens SECIS binding protein 2 (SBP2), mRNA
NM_022730	Homo sapiens COP9 constitutive photomorphogenic homolog subunit 7B
	(Arabidopsis) (COPS7B), mRNA
NM_022574	Homo sapiens postmeiotic segregation increased 2-like 12 (PERQ1), mRNA
NM_022568	Homo sapiens aldehyde dehyrdogenase 8 family, member A1 (ALDH8A1), mRNA
NM_022477	Homo sapiens NDRG family member 3 (NDRG3), mRNA
NM_022082	Homo sapiens chromosome 20 open reading frame 59 (C20orf59), mRNA
NM_022058	Homo sapiens solute carrier family 4, sodium bicarbonate transporter-like,
	member 10 (SLC4A10), mRNA
NM_021230	Homo sapiens myeloid/lymphoid or mixed-lineage leukemia3 (MLL3), mRNA
NM_021145	Homo sapiens cyclin D binding myb-like transcription factor 1 (DMTF1), mRNA
NM_005238	Homo sapiens v-ets erythroblastosis virus E26 oncogene homolog 1 (avian)
	(ETS1), mRNA
NM_020465	Homo sapiens NDRG family member 4 (NDRG4), mRNA
NM_014227	Homo sapiens solute carrier family 5 (low affinity glucose cotransporter), member 4 (SLC5A4), mRNA
NM 015317	Homo sapiens pumilio homolog 2 (Drosophila) (PUM2), mRNA
NM_015665	Homo sapiens achalasia, adrenocortical insufficiency, alacrimia (Allgrove, triple-A) (AAAS), mRNA
NM 021950	Homo sapiens membrane-spanning 4-domains, subfamily A, member 2 (Fc
111/2_0225	fragment of IgE, high affinity I, receptor for; beta polypeptide) (MS4A1), mRNA
NM_005589	Homo sapiens aldehyde dehydrogenase 6 family, member A1 (ALDH6A1), mRNA
NM 000533	Homo sapiens proteolipid protein1 (Pelizaeus-Merzbacher disease, spastic
	paraplegia 2, uncomplicated) (PLP1), mRNA
NM 016252	Homo sapiens baculoviral IAP repeat-containing 6 (apollon) (BIRC6), mRNA
NM 014351	Homo sapiens sulfotransferase family 4A, member 1 (SULT4A1), mRNA
NM_012323	Homo sapiens v-maf musculoaponeurotic fibrosarcoma oncogene homolog F (avian) (MAFF), mRNA
NM_006600	Homo sapiens nuclear distribution gene C homolog (A. nidulans) (NUDC), mRNA
NM_006145	Homo sapiens DnaJ (Hsp40) homolog, subfmaily B, member 1 (DNAJB1), mRNA
NM_005120	Homo sapiens trinucleotide repeat containing 11 (THR-associated protein, 230 kD subunit) (TNRC11), mRNA
NM_001383	Homo sapiens diptheria toxin resistance protein required for diphthamide
_	biosynthesis-like 1 (S. cerevisiae) (DPH2L1), mRNA
NM 001327	Homo sapiens cancer/testis antigen 1 (CTAG1), mRNA
NM 080750	Homo sapiens chromosome 20 open reading frame 143 (C20orf143), mRNA
NM 032819	Homo sapiens zinc finger protein 341 (ZNF341), mRNA
NM_017895	Homo sapiens DEAD/H (Asp-Glu-Ala-Asp/His) box polypeptide 27 (DDX27), mRNA
NM 030782	Homo sapiens cisplatin resistance related protein CRR9p (CRR9), mRNA
NM 080748	Homo sapiens chromosome 20 open reading frame 52 (C20orf52), mRNA
NM 080743	Homo sapiens serine-arginine repressor protein (35 kDa) (SRrp35), mRNA
NM 080742	Homo sapiens UDP-glucuronyltransferase-S (GLCATS), mRNA

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COX7A2 , nuclear gene encoding mitochondrial protein, mRNA		
NM_001864 Homo sapiens cytochrome c oxidase subunit VIIa polypeptide 1 (muscle) (COX7AI), nuclear gene encoding mitochondrial protein, mRNA NM_0060544 Homo sapiens collectin sub-family member 10 (C-type lectin) (COLEC10), mRNA NM_080544 Homo sapiens collagen-like tail subunit (single strand of homotrimer) of asymmetric acetylcholinesterase (COLQ), transcript variant VIII, mRNA NM_080543 Homo sapiens collagen-like tail subunit (single strand of homotrimer) of asymmetric acetylcholinesterase (COLQ), transcript variant VI, mRNA NM_080542 Homo sapiens collagen-like tail subunit (single strand of homotrimer) of asymmetric acetylcholinesterase (COLQ), transcript variant VI, mRNA NM_080541 Homo sapiens collagen-like tail subunit (single strand of homotrimer) of asymmetric acetylcholinesterase (COLQ), transcript variant V, mRNA NM_080540 Homo sapiens collagen-like tail subunit (single strand of homotrimer) of asymmetric acetylcholinesterase (COLQ), transcript variant IV, mRNA NM_080539 Homo sapiens collagen-like tail subunit (single strand of homotrimer) of asymmetric acetylcholinesterase (COLQ), transcript variant II, mRNA NM_080534 Homo sapiens collagen-like tail subunit (single strand of homotrimer) of asymmetric acetylcholinesterase (COLQ), transcript variant II, mRNA NM_080535 Homo sapiens sapotosis related protein APR-3 (APR-3), transcript variant 2, mRNA NM_080540 Homo sapiens apoptosis related protein APR-3 (APR-3), transcript variant 2, mRNA NM_080550 Homo sapiens apoptosis related protein APR-3 (APR-3), transcript variant 1, mRNA NM_080540 Homo sapiens sapotosis related protein (APR-2), mRNA NM_080541 Homo sapiens sadaptor-related protein complex 1, gamma 1 subunit (APIGI), mRNA NM_080545 Homo sapiens adaptor-related protein complex 1, gamma 2 subunit (APIGI), mRNA NM_080545 Homo sapiens adaptor-related protein complex 1, gamma 2 subunit (APIGI), transcript variant 1, mRNA NM_080546 Homo sapiens protein tyrosine phosphatase, non-receptor type 6 (PTPN6), transcript variant 1, mRNA NM_080548 Homo sapiens protei	NM_001865	Homo sapiens cytochrome c oxidase subunit VIIa polypeptide 2 (liver)
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NM_080542 Homo sapiens collagen-like tail subunit (single strand of homotrimer) of asymmetric acetylcholinesterase (COLQ), transcript variant VI, mRNA	1111_000515	asymmetric acetylcholinesterase (COLQ), transcript variant VII, mRNA
asymmetric acetylcholinesterase (COLQ), transcript variant VI, mRNA NM_080540 Homo sapiens collagen-like tail subunit (single strand of homotrimer) of asymmetric acetylcholinesterase (COLQ), transcript variant V, mRNA NM_080540 Homo sapiens collagen-like tail subunit (single strand of homotrimer) of asymmetric acetylcholinesterase (COLQ), transcript variant IV, mRNA NM_080538 Homo sapiens collagen-like tail subunit (single strand of homotrimer) of asymmetric acetylcholinesterase (COLQ), transcript variant III, mRNA NM_080538 Homo sapiens collagen-like tail subunit (single strand of homotrimer) of asymmetric acetylcholinesterase (COLQ), transcript variant II, mRNA NM_080567 Homo sapiens collagen-like tail subunit (single strand of homotrimer) of asymmetric acetylcholinesterase (COLQ), transcript variant II, mRNA NM_080592 Homo sapiens apoptosis related protein APR-3 (APR-3), transcript variant 2, mRNA NM_016085 Homo sapiens apoptosis related protein APR-3 (APR-3), transcript variant 1, mRNA NM_0101745 Homo sapiens apoptosis related protein (APR-2), mRNA NM_0014311 Homo sapiens calcium modulating ligand (CAMLG), mRNA NM_004341 Homo sapiens calcium modulating ligand (CAMLG), mRNA NM_004341 NM_0032493 Homo sapiens adaptor-related protein complex 1, mul 1 subunit (APIM1), mRNA NM_001128 Homo sapiens adaptor-related protein complex 1, gamma 1 subunit (APIG1), mRNA NM_0080545 Homo sapiens adaptor-related protein complex 1, gamma 2 subunit (APIG2), transcript variant 1, mRNA NM_080549 Homo sapiens adaptor-related protein complex 1, gamma 2 subunit (APIG2), transcript variant 1, mRNA NM_080549 Homo sapiens protein tyrosine phosphatase, non-receptor type 6 (PTPN6), transcript variant 1, mRNA NM_080549 Homo sapiens protein tyrosine phosphatase, non-receptor type 6 (PTPN6), transcript variant 1, mRNA NM_002830 Homo sapiens protein tyrosine phosphatase, non-receptor type 6 (PTPN6), transcript variant 1, mRNA NM_00289 Homo sapiens protein tyrosine phosphatase, non-receptor type 6 (PTPN6), transcript variant	NIM 080542	Homo saniens collagen-like tail subunit (single strand of homotrimer) of
NM_080541 Homo sapiens collagen-like tail subunit (single strand of homotrimer) of asymmetric acetylcholinesterase (COLQ), transcript variant V, mRNA	14141_000542	asymmetric acetylcholinesterase (COLO), transcript variant VI, mRNA
asymmetric acetylcholinesterase (COLQ), transcript variant V, mRNA NM_080530 Homo sapiens collagen-like tail subunit (single strand of homotrimer) of asymmetric acetylcholinesterase (COLQ), transcript variant IV, mRNA NM_080538 Homo sapiens collagen-like tail subunit (single strand of homotrimer) of asymmetric acetylcholinesterase (COLQ), transcript variant III, mRNA NM_080538 Homo sapiens collagen-like tail subunit (single strand of homotrimer) of asymmetric acetylcholinesterase (COLQ), transcript variant III, mRNA NM_005677 Homo sapiens collagen-like tail subunit (single strand of homotrimer) of asymmetric acetylcholinesterase (COLQ), transcript variant I, mRNA NM_080592 Homo sapiens apoptosis related protein APR-3 (APR-3), transcript variant 2, mRNA NM_016085 Homo sapiens apoptosis related protein APR-3 (APR-3), transcript variant 1, mRNA NM_0104318 Homo sapiens apoptosis related protein (APR-2), mRNA NM_001745 Homo sapiens apoptosis related protein (APR-2), mRNA NM_004341 Homo sapiens carbamoyl-phosphate synthetase 2, aspartate transcarbamylase, and dihydrorotase (CAD),, nuclear gene encoding mitochondrial protein, mRNA NM_001128 Homo sapiens adaptor-related protein complex 1, mu 1 subunit (APIM1), mRNA NM_001128 Homo sapiens adaptor-related protein complex 1, gamma 2 subunit (APIG1), mRNA NM_003917 Homo sapiens adaptor-related protein complex 1, gamma 2 subunit (APIG2), transcript variant 1, mRNA NM_080549 Homo sapiens protein tyrosine phosphatase, non-receptor type 6 (PTPN6), transcript variant 1, mRNA NM_080549 Homo sapiens protein tyrosine phosphatase, non-receptor type 6 (PTPN6), transcript variant 1, mRNA NM_002831 Homo sapiens protein tyrosine phosphatase, non-receptor type 6 (PTPN6), transcript variant 1, mRNA NM_002830 Homo sapiens protein tyrosine phosphatase, non-receptor type 6 (PTPN6), transcript variant 1, mRNA NM_002830 Homo sapiens protein tyrosine phosphatase, non-receptor type 6 (PTPN6), transcript variant 3, mRNA	NIM 080541	Homo saniens collagen-like tail subunit (single strand of homotrimer) of
NM_080540	14101_000341	asymmetric acetylcholinesterase (COLO) transcript variant V. mRNA
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transcript variant 1, mRNA NM_080549 Homo sapiens protein tyrosine phosphatase, non-receptor type 6 (PTPN6), transcript variant 3, mRNA NM_080548 Homo sapiens protein tyrosine phosphatase, non-receptor type 6 (PTPN6), transcript variant 2, mRNA NM_002831 Homo sapiens protein tyrosine phosphatase, non-receptor type 6 (PTPN6), transcript variant 1, mRNA NM_002830 Homo sapiens protein tyrosine phosphatase, non-receptor type 4 (megakaryocyte) (PTPN4), mRNA NM_002829 Homo sapiens protein tyrosine phosphatase, non-receptor type 3 (PTPN3), mRNA NM_080423 Homo sapiens protein tyrosine phosphatase, non-receptor type 2 (PTPN2), transcript variant 3, mRNA	NIM 003017	Homo saniens adaptor-related protein complex 1, gamma 2 subunit (AP1G2).
NM_080549 Homo sapiens protein tyrosine phosphatase, non-receptor type 6 (PTPN6), transcript variant 3, mRNA NM_080548 Homo sapiens protein tyrosine phosphatase, non-receptor type 6 (PTPN6), transcript variant 2, mRNA NM_002831 Homo sapiens protein tyrosine phosphatase, non-receptor type 6 (PTPN6), transcript variant 1, mRNA NM_002830 Homo sapiens protein tyrosine phosphatase, non-receptor type 4 (megakaryocyte) (PTPN4), mRNA NM_002829 Homo sapiens protein tyrosine phosphatase, non-receptor type 3 (PTPN3), mRNA NM_080423 Homo sapiens protein tyrosine phosphatase, non-receptor type 2 (PTPN2), transcript variant 3, mRNA	14141_003717	
transcript variant 3, mRNA NM_080548 Homo sapiens protein tyrosine phosphatase, non-receptor type 6 (PTPN6), transcript variant 2, mRNA NM_002831 Homo sapiens protein tyrosine phosphatase, non-receptor type 6 (PTPN6), transcript variant 1, mRNA NM_002830 Homo sapiens protein tyrosine phosphatase, non-receptor type 4 (megakaryocyte) (PTPN4), mRNA NM_002829 Homo sapiens protein tyrosine phosphatase, non-receptor type 3 (PTPN3), mRNA NM_080423 Homo sapiens protein tyrosine phosphatase, non-receptor type 2 (PTPN2), transcript variant 3, mRNA	NIM DROSAD	
NM_080548 Homo sapiens protein tyrosine phosphatase, non-receptor type 6 (PTPN6), transcript variant 2, mRNA NM_002831 Homo sapiens protein tyrosine phosphatase, non-receptor type 6 (PTPN6), transcript variant 1, mRNA NM_002830 Homo sapiens protein tyrosine phosphatase, non-receptor type 4 (megakaryocyte) (PTPN4), mRNA NM_002829 Homo sapiens protein tyrosine phosphatase, non-receptor type 3 (PTPN3), mRNA NM_080423 Homo sapiens protein tyrosine phosphatase, non-receptor type 2 (PTPN2), transcript variant 3, mRNA	14141 000243	
transcript variant 2, mRNA NM_002831 Homo sapiens protein tyrosine phosphatase, non-receptor type 6 (PTPN6), transcript variant 1, mRNA NM_002830 Homo sapiens protein tyrosine phosphatase, non-receptor type 4 (megakaryocyte) (PTPN4), mRNA NM_002829 Homo sapiens protein tyrosine phosphatase, non-receptor type 3 (PTPN3), mRNA NM_080423 Homo sapiens protein tyrosine phosphatase, non-receptor type 2 (PTPN2), transcript variant 3, mRNA	ND4 000540	
NM_002831 Homo sapiens protein tyrosine phosphatase, non-receptor type 6 (PTPN6), transcript variant 1, mRNA NM_002830 Homo sapiens protein tyrosine phosphatase, non-receptor type 4 (megakaryocyte) (PTPN4), mRNA NM_002829 Homo sapiens protein tyrosine phosphatase, non-receptor type 3 (PTPN3), mRNA NM_080423 Homo sapiens protein tyrosine phosphatase, non-receptor type 2 (PTPN2), transcript variant 3, mRNA	NMI_U8U348	
transcript variant 1, mRNA NM_002830 Homo sapiens protein tyrosine phosphatase, non-receptor type 4 (megakaryocyte) (PTPN4), mRNA NM_002829 Homo sapiens protein tyrosine phosphatase, non-receptor type 3 (PTPN3), mRNA NM_080423 Homo sapiens protein tyrosine phosphatase, non-receptor type 2 (PTPN2), transcript variant 3, mRNA	2D (00000:	
NM_002830 Homo sapiens protein tyrosine phosphatase, non-receptor type 4 (megakaryocyte) (PTPN4), mRNA NM_002829 Homo sapiens protein tyrosine phosphatase, non-receptor type 3 (PTPN3), mRNA NM_080423 Homo sapiens protein tyrosine phosphatase, non-receptor type 2 (PTPN2), transcript variant 3, mRNA	NM_002831	
(megakaryocyte) (PTPN4), mRNA NM_002829 Homo sapiens protein tyrosine phosphatase, non-receptor type 3 (PTPN3), mRNA NM_080423 Homo sapiens protein tyrosine phosphatase, non-receptor type 2 (PTPN2), transcript variant 3, mRNA	200000	
NM_002829 Homo sapiens protein tyrosine phosphatase, non-receptor type 3 (PTPN3), mRNA NM_080423 Homo sapiens protein tyrosine phosphatase, non-receptor type 2 (PTPN2), transcript variant 3, mRNA	NM_002830	
mRNA NM_080423 Homo sapiens protein tyrosine phosphatase, non-receptor type 2 (PTPN2), transcript variant 3, mRNA		(megakaryocyte) (P1PN4), mKNA
transcript variant 3, mRNA	NM_002829	mRNA
	NM_080423	
	NM 080422	Homo sapiens protein tyrosine phosphatase, non-receptor type 2 (PTPN2),

	transcript variant 2, mRNA
NM_002828	Homo sapiens protein tyrosine phosphatase, non-receptor type 2 (PTPN2),
	transcript variant 1, mRNA
NM_002827	Homo sapiens protein tyrosine phosphatase, non-receptor type 1 (PTPN1), mRNA
NM_014241	Homo sapiens protein tyrosine phosphatase-like (proline instead of catalytic arginine), member a (PTPLA), mRNA
NM_003479	Homo sapiens protein tyrosine phosphatase type IVA, member 2 (PTP4A2), transcript variant 1, mRNA
NM_080392	Homo sapiens protein tyrosine phosphatase type IVA, member 2 (PTP4A2), transcript variant 3, mRNA
NM_080391	Homo sapiens protein tyrosine phosphatase type IVA, member 2 (PTP4A2), transcript variant 2, mRNA
NM_080591	Homo sapiens prostaglandin-endoperoxide synthase 1 (prostaglandin G/H synthase and cyclooxygenase) (PTGS1), transcript variant 2, mRNA
NM_000962	Homo sapiens prostaglandin-endoperoxide synthase 1 (prostaglandin G/H synthase and cyclooxygenase) (PTGS1), transcript variant 1, mRNA
NM 004058	Homo sapiens calcyphosine (CAPS), transcript variant 1, mRNA
NM 080590	Homo sapiens calcyphosine (CAPS), transcript variant 2, mRNA
NM_006380	Homo sapiens amyloid beta precursor protein (cytoplasmic tail) binding protein 2 (APPBP2), mRNA
NM_003905	Homo sapiens amyloid beta precursor protein binding protein 1, 59kD (APPBP1), mRNA
NM_005783	Homo sapiens ATP binding protein associated with cell differentiation (APACD), mRNA
NM_080600	Homo sapiens myelin associated glycoprotein (MAG), transcript variant 2, mRNA
NM_002361	Homo sapiens myelin associated glycoprotein (MAG), transcript variant 1, mRNA
NM 005994	Homo sapiens T-box 2 (TBX2), mRNA
NM 080647	Homo sapiens T-box 1 (TBX1), transcript variant C, mRNA
NM 080646	Homo sapiens T-box 1 (TBX1), transcript variant A, mRNA
NM 080675	Homo sapiens sperm associated antigen 4-like (SPAG4L), mRNA
NM 080617	Homo sapiens cerebellin precursor-like 1 (CBLNL1), mRNA
NM 080611	Homo sapiens dual specificity phosphatase-like 15 (DUSP15), mRNA
NM 080610	Homo sapiens cystatin 9-like (mouse) (CST9L), mRNA
NM_080602	Homo sapiens actin related protein 2/3 complex, subunit 3B (21 kD) (ARPC3B), mRNA
NG_000011	Homo sapiens genomic cytochrome P450, subfamily IIA (phenobarbital-inducible) (CYP2A.3@) on chromosome 19
NM_016649	Homo sapiens chromosome 20 open reading frame 6 (C20orf6), mRNA
NM_080597	Homo sapiens oxysterol binding protein-like 1A (OSBPL1A), mRNA
NM_080605	Homo sapiens UDP-Gal:betaGlcNAc beta 1,3-galactosyltransferase, polypeptide 6 (B3GALT6), mRNA
NM_058169	Homo sapiens loss of heterozygosity, 12, chromosomal region 1 (LOH12CR1), mRNA
NM_058164	Homo sapiens olfactomedin 2 (OLFM2), mRNA
NM 052866	Homo sapiens ADAMTS-like 1 (ADAMTSL1), mRNA
NM 018030	Homo sapiens oxysterol binding protein-like 1A (OSBPL1A), mRNA
NM 033142	Homo sapiens chorionic gonadotropin, beta polypeptide 7 (CGB7), mRNA
NG_000013	Homo sapiens genomic MHC class III complement gene cluster (MCGC@) on chromosome 6

NM_020967	Homo sapiens nuclear receptor coactivator 5 (NCOA5), mRNA
NM_033044	Homo sapiens microtubule-actin crosslinking factor 1 (MACF1), transcript
	variant 3, mRNA
NM 033024	Homo sapiens microtubule-actin crosslinking factor 1 (MACF1), transcript
_	variant 2, mRNA
NG 000017	Homo sapiens genomic protocadherin beta cluster (PCDHB@) on chromosome 5
NM 015864	Homo sapiens chromosome 6 open reading frame 32 (C6orf32), mRNA
NM 032188	Homo sapiens histone acetyltransferase MYST1 (MYST1), mRNA
NM 030776	Homo sapiens chromosome 20 open reading frame 183 (C20orf183), mRNA
NM 024918	Homo sapiens chromosome 20 open reading frame 172 (C20orf172), mRNA
NM 024812	Homo sapiens brain and acute leukemia, cytoplasmic (BAALC), mRNA
NM 024777	Homo sapiens chromosome 20 open reading frame 124 (C20orf124), mRNA
NM 024758	Homo sapiens agmatinase (FLJ23384), mRNA
NM 024641	Homo sapiens mandaselin (FLJ12838), mRNA
NM 024331	Homo sapiens chromosome 20 open reading frame 121 (C20orf121), mRNA
NM 024301	Homo sapiens fukutin-related protein (FKRP), mRNA
NM 005763	Homo sapiens aminoadipate-semialdehyde synthase (AASS), mRNA
NM 023935	Homo sapiens chromosome 20 open reading frame 116 (C20orf116), mRNA
NM_021993	Homo sapiens FUS interacting protein (serine-arginine rich) 2 (FUSIP2), mRNA
NM 014555	Homo sapiens transient receptor potential cation channel, subfamily M, member
14141_014555	5 (TRPM5), mRNA
NM 000537	Homo sapiens renin (REN), mRNA
NM_016652	Homo sapiens Crn, crooked neck-like 1 (Drosophila) (CRNKL1), mRNA
NM 021245	Homo sapiens myozenin 1 (MYOZ1), mRNA
NM 001967	Homo sapiens eukaryotic translation initiation factor 4A, isoform 2 (EIF4A2),
14141_001507	mRNA
NM 018649	Homo sapiens H2A histone family, member Y2 (H2AFY2), mRNA
NM 015148	Homo sapiens PAS domain containing serine/threonine kinase (PASK), mRNA
NM 017902	Homo sapiens hypoxia-inducible factor 1, alpha subunit inhibitor (HIF1AN),
11112017702	mRNA
NM 018285	Homo sapiens chromosome 15 open reading frame 12 (C15orf12), nuclear gene
11172_010200	encoding mitochondrial protein, mRNA
NM 018267	Homo sapiens H2A histone family, member J (H2AFJ), mRNA
NM 017555	Homo sapiens egl nine homolog 2 (C. elegans) (EGLN2), transcript variant 2,
11112017555	mRNA
NM 016143	Homo sapiens likely ortholog of rat p47 (p47), mRNA
NM 015993	Homo sapiens plasmolipin (PMLP), mRNA
NM 014938	Homo sapiens Mlx interactor (MONDOA), mRNA
NM_014948	Homo sapiens likely ortholog of mouse ubiquitin conjugating enzyme 7
1111_017510	interacting protein 5 (UBCE7IP5), mRNA
NM 014016	Homo sapiens SAC1 suppressor of actin mutations 1-like (yeast) (SACM1L),
1111_014010	mRNA
NM 015156	Homo sapiens REST corepressor (RCOR), mRNA
NM 013337	Homo sapiens translocase of inner mitochondrial membrane 22 homolog (yeast)
14112013337	(TIMM22), mRNA
NM 013233	Homo sapiens serine threonine kinase 39 (STE20/SPS1 homolog, yeast)
1414_015255	(STK39), mRNA
NM 006595	Homo sapiens apoptosis inhibitor 5 (API5), mRNA
NM 006402	Homo sapiens hepatitis B virus x interacting protein (HBXIP), mRNA
NM 006351	Homo sapiens translocase of inner mitochondrial membrane 44 homolog (yeast)
14147_000221	(TIMM44), mRNA
NM_006327	Homo sapiens translocase of inner mitochondrial membrane 23 homolog (yeast)
1111 000001	1

	(TIMM23), mRNA
NM_006335	Homo sapiens translocase of inner mitochondrial membrane 17 homolog A
141/41 _000222	(yeast) (TIMM17A), mRNA
NM_006420	Homo sapiens ADP-ribosylation factor guanine nucleotide-exchange factor 2
14141_000420	(brefeldin A-inhibited) (ARFGEF2), mRNA
NM 005992	Homo sapiens T-box 1 (TBX1), transcript variant B, mRNA
	Homo sapiens translocase of inner mitochondrial membrane 17 homolog B
NM_005834	(yeast) (TIMM17B), mRNA
NM_000385	Homo sapiens aquaporin 1 (channel-forming integral protein, 28kD) (AQP1), mRNA
NM_002891	Homo sapiens Ras protein-specific guanine nucleotide-releasing factor 1 (RASGRF1), mRNA
NM 000963	Homo sapiens prostaglandin-endoperoxide synthase 2 (prostaglandin G/H
_	synthase and cyclooxygenase) (PTGS2), mRNA
NM_002792	Homo sapiens proteasome (prosome, macropain) subunit, alpha type, 7 (PSMA7), mRNA
NM 002335	Homo sapiens low density lipoprotein receptor-related protein 5 (LRP5), mRNA
NM_001402	Homo sapiens eukaryotic translation elongation factor 1 alpha 1 (EEF1A1), mRNA
NM 080677	Homo sapiens dynein light chain 2 (Dlc2), mRNA
NM 080672	Homo sapiens Q9H4T4 like (H17739), mRNA
NM 080671	Homo sapiens potassium voltage-gated channel, Isk-related subfamily, gene 4
944	(KCNE4), mRNA
NM 080670	Homo sapiens similar to RIKEN cDNA 2610030J16 gene (MGC2541), mRNA
NM 080669	Homo sapiens similar to RIKEN cDNA 1110002C08 gene (MGC9564), mRNA
NM 080667	Homo sapiens similar to RIKEN cDNA 4931428D14 gene (MGC15407), mRNA
NM_080665	Homo sapiens similar to RIKEN cDNA B230118G17 gene (MGC19604), mRNA
NM 080664	Homo sapiens similar to RIKEN cDNA 4930578F06 gene (MGC9912), mRNA
NM 080662	Homo sapiens similar to RIKEN cDNA 1810022F11 gene (MGC4281), mRNA
NM 080660	Homo sapiens similar to RIKEN cDNA 1200014N16 gene (MGC14289), mRNA
NM_080659	Homo sapiens similar to RIKEN cDNA 2310030G06 gene (MGC14839), mRNA
NM 080657	Homo sapiens vipirin (cig5), mRNA
NM 080655	Homo sapiens similar to RIKEN cDNA 5730528L13 gene (MGC17337), mRNA
NM 080654	Homo sapiens NY-REN-41 antigen (NY-REN-41), mRNA
NM 080653	Homo sapiens similar to RIKEN cDNA 4930500C14 gene (MGC9341), mRNA
NM 080652	Homo sapiens similar to RIKEN cDNA 5730578N08 gene (MGC15397), mRNA
NM 004296	Homo sapiens regulator of G-protein signalling 6 (RGS6), mRNA
NM 014234	Homo sapiens FabG (beta-ketoacyl-[acyl-carrier-protein] reductase, E coli) like
1111_01-25	(E. coli) (FABGL), mRNA
NM 024775	Homo sapiens gemin 6 (GEMIN6), mRNA
NM_080626	Homo sapiens BRI3 binding protein (BRI3BP), mRNA
NM 080625	Homo sapiens chromosome 20 open reading frame 160 (C20orf160), mRNA
NM_080616	Homo sapiens chromosome 20 open reading frame 112 (C20orf112), mRNA
NM 080612	Homo sapiens DOS/Gab family member 3 (GAB3), mRNA
NM 080607	Homo sapiens bos/dab family inclined 5 (GAB5), inktyA Homo sapiens chromosome 20 open reading frame 102 (C20orf102), mRNA
NM_080603	Homo sapiens chromosome 20 open reading frame 162 (C20orf162), mRNA
NM_032019	Homo sapiens chromosome 20 open reading frame 102 (C2001102), firkNA Homo sapiens histone deacetylase 10 (HDAC10), mRNA
NM 030815	Homo sapiens chromosome 20 open reading frame 126 (C20orf126), mRNA
NM 020841	Homo sapiens enromosome 20 open reading frame 126 (C2001126), InRNA Homo sapiens oxysterol binding protein-like 8 (OSBPL8), mRNA
NM 020764	Homo sapiens oxysteror briding protein-like 8 (OSBFL8), INKINA Homo sapiens cask-interacting protein 1 (CASKIN1), mRNA
NM_016436	Homo sapiens cask-interacting protein 1 (CASKIN1), mRNA Homo sapiens chromosome 20 open reading frame 104 (C20orf104), mRNA
14141 010430	1 1101110 Sapiens chromosome 20 open reading frame 104 (C20011104), mRNA

NM_022104	Homo sapiens chromosome 20 open reading frame 67 (C20orf67), mRNA
NM_080546	Homo sapiens CDw92 antigen (CDW92), mRNA
NM_015511	Homo sapiens chromosome 20 open reading frame 4 (C20orf4), mRNA
NM 002116	Homo sapiens major histocompatibility complex, class I, A (HLA-A), mRNA
NM_023017	Homo sapiens phosphoinositide 3-kinase enhancer (PIKE), mRNA
NM 020933	Homo sapiens zinc finger protein 317 (ZNF317), mRNA
NM_005037	Homo sapiens peroxisome proliferative activated receptor, gamma (PPARG), mRNA
NM 018206	Homo sapiens vacuolar protein sorting 35 (yeast) (VPS35), mRNA
NM_014003	Homo sapiens DEAD/H (Asp-Glu-Ala-Asp/His) box polypeptide 38 (DDX38), mRNA
NM_006445	Homo sapiens PRP8 pre-mRNA processing factor 8 homolog (yeast) (PRPF8), mRNA
NM_003675	Homo sapiens pre-mRNA processing factor 18 (PRP18), mRNA
NM_006214	Homo sapiens phytanoyl-CoA hydroxylase (Refsum disease) (PHYH), mRNA
NM_004374	Homo sapiens cytochrome c oxidase subunit VIc (COX6C), nuclear gene encoding mitochondrial protein, mRNA
NM_001863	Homo sapiens cytochrome c oxidase subunit VIb (COX6B), nuclear gene encoding mitochondrial protein, mRNA
NM_005205	Homo sapiens cytochrome c oxidase subunit VIa polypeptide 2 (COX6A2), nuclear gene encoding mitochondrial protein, mRNA
NM_004373	Homo sapiens cytochrome c oxidase subunit VIa polypeptide 1 (COX6A1), nuclear gene encoding mitochondrial protein, mRNA
NM_032609	Homo sapiens cytochrome c oxidase subunit IV isoform 2 (COX4I2), nuclear gene encoding mitochondrial protein, mRNA
NM 032489	Homo sapiens acrosin binding protein (ACRBP), mRNA
NM_080476	Homo sapiens CDC91 cell division cycle 91-like 1 (S. cerevisiae) (CDC91L1), mRNA
NM_080473	Homo sapiens GATA binding protein 5 (GATA5), mRNA
NM_002121	Homo sapiens major histocompatibility complex, class II, DP beta 1 (HLA-DPB1), mRNA
NM_078470	Homo sapiens COX15 homolog, cytochrome c oxidase assembly protein (yeast) (COX15), nuclear gene encoding mitochondrial protein, transcript variant 1, mRNA
NM_004375	Homo sapiens COX11 homolog, cytochrome c oxidase assembly protein (yeast) (COX11), nuclear gene encoding mitochondrial protein, mRNA
NM_001303	Homo sapiens COX10 homolog, cytochrome c oxidase assembly protein, heme A/farnesyltransferase (yeast) (COX10), nuclear gene encoding mitochondrial protein, mRNA
NM_054028	Homo sapiens acyl-malonyl condensing enzyme (AMAC), mRNA
NM_032485	Homo sapiens chromosome 20 open reading frame 154 (C20orf154), mRNA
NM_033342	Homo sapiens tripartite motif-containing 7 (TRIM7), mRNA
NM_033421	Homo sapiens chromosome 20 open reading frame 161 (C20orf161), mRNA
NM_033197	Homo sapiens chromosome 20 open reading frame 114 (C20orf114), mRNA
NM_020866	Homo sapiens kelch-like 1 (Drosophila) (KLHL1), mRNA
NM_032883	Homo sapiens chromosome 20 open reading frame 100 (C20orf100), mRNA
NM_032523	Homo sapiens oxysterol binding protein-like 6 (OSBPL6), mRNA
NM_020896	Homo sapiens oxysterol binding protein-like 5 (OSBPL5), mRNA
NM_015550	Homo sapiens oxysterol binding protein-like 3 (OSBPL3), mRNA
NM_031473	Homo sapiens carnitine deficiency-associated gene expressed in ventricle 1 (CDV-1), mRNA
NM_030801	Homo sapiens MAGE-E1 protein (MAGE-E1), mRNA

NIM 025129	Home seniors MISS1 endenuelesse (MISS1) mDNA
NM_025128	Homo sapiens MUS81 endonuclease (MUS81), mRNA Homo sapiens altromosome 20 open reading frame 98 (C20orf08) mRNA
NM_024958 NM_024663	Homo sapiens chromosome 20 open reading frame 98 (C20orf98), mRNA
	Homo sapiens aminopeptidase-like 1 (NPEPL1), mRNA
NM_024586	Homo sapiens oxysterol binding protein-like 9 (OSBPL9), mRNA
NM_024120	Homo sapiens chromosome 20 open reading frame 7 (C20orf7), mRNA
NM_022776	Homo sapiens oxysterol binding protein-like 11 (OSBPL11), mRNA
NM_022109	Homo sapiens CDw92 antigen (CDW92), mRNA
NM_022088	Homo sapiens zinc finger protein 338 (ZNF338), mRNA
NM_021158	Homo sapiens chromosome 20 open reading frame 97 (C20orf97), mRNA
NM_021232	Homo sapiens proline dehydrogenase (oxidase) 2 (PRODH2), mRNA
NM_021220	Homo sapiens zinc finger protein 339 (ZNF339), mRNA
NM_021039	Homo sapiens S100 calcium binding protein A14 (calgizzarin) (S100A14), mRNA
NM_020659	Homo sapiens tweety homolog 1 (Drosophila) (TTYH1), mRNA
NM_018972	Homo sapiens ganglioside-induced differentiation-associated protein 1 (GDAP1), mRNA
NM 017921	Homo sapiens hypothetical protein FLJ20657 (NPL4), mRNA
NM_017784	Homo sapiens oxysterol binding protein-like 10 (OSBPL10), mRNA
NM 017731	Homo sapiens oxysterol binding protein-like 7 (OSBPL7), mRNA
NM_018209	Homo sapiens ADP-ribosylation factor 1 GTPase activating protein (ARF1GAP), mRNA
NM 018102	Homo sapiens zinc finger protein 334 (ZNF334), mRNA
NM 015891	Homo sapiens pre-mRNA splicing factor 17 (PRP17), mRNA
NM 016599	Homo sapiens myozenin 2 (MYOZ2), mRNA
NM 014962	Homo sapiens BTB (POZ) domain containing 3 (BTBD3), mRNA
NM 014835	Homo sapiens oxysterol binding protein-like 2 (OSBPL2), mRNA
NM 014723	Homo sapiens syntaphilin (SNPH), mRNA
NM 014183	Homo sapiens dynein light chain 2A (DNLC2A), mRNA
NM_014055	Homo sapiens carnitine deficiency-associated gene expressed in ventricle 1 (CDV-1), mRNA
NM 014477	Homo sapiens chromosome 20 open reading frame 10 (C20orf10), mRNA
NM 012261	Homo sapiens chromosome 20 open reading frame 103 (C20orf103), mRNA
NM 013369	Homo sapiens DNA (cytosine-5-)-methyltransferase 3-like (DNMT3L), mRNA
NM 012469	Homo sapiens chromosome 20 open reading frame 14 (C20orf14), mRNA
NM 012291	Homo sapiens extra spindle poles like 1 (S. cerevisiae) (ESPL1), mRNA
NM 007002	Homo sapiens adhesion regulating molecule 1 (ADRM1), mRNA
NM_006809	Homo sapiens translocase of outer mitochondrial membrane 34 (TOMM34), mRNA
NM 006813	Homo sapiens proline rich 2 (PROL2), mRNA
NM_002509	Homo sapiens NK2 transcription factor homolog B (Drosophila) (NKX2B), mRNA
NM_080474	Homo sapiens serine (or cysteine) proteinase inhibitor, clade B (ovalbumin), member 12 (SERPINB12), mRNA
NM 006009	Homo sapiens tubulin, alpha 3 (TUBA3), mRNA
NM_003463	Homo sapiens protein tyrosine phosphatase type IVA, member 1 (PTP4A1), mRNA
NM 019888	Homo sapiens melanocortin 3 receptor (MC3R), mRNA
NM 001846	Homo sapiens collagen, type IV, alpha 2 (COLAA2), mRNA
NM_079422	Homo sapiens myosin, light polypeptide 1, alkali; skeletal, fast (MYL1), transcript variant 3f, mRNA
NM_079420	Homo sapiens myosin, light polypeptide 1, alkali; skeletal, fast (MYL1), transcript variant 1f, mRNA

NM_00795 Homo sapiens dopamine receptor D2 (DRD2), transcript variant 1, mRNA NM_016574 Homo sapiens dopamine receptor D2 (DRD2), transcript variant 2, mRNA NM_079837 Homo sapiens BTG3 associated nuclear protein (BANP), transcript variant 2, mRNA NM_017869 Homo sapiens BTG3 associated nuclear protein (BANP), transcript variant 1, mRNA NM_079425 Homo sapiens myosin, light polypeptide 6, alkali, smooth muscle and non- muscle (MYL6), transcript variant 3, mRNA NM_079424 Homo sapiens myosin, light polypeptide 6, alkali, smooth muscle and non- muscle (MYL6), transcript variant 4, mRNA NM_079423 Homo sapiens myosin, light polypeptide 6, alkali, smooth muscle and non- muscle (MYL6), transcript variant 2, mRNA NM_021019 Homo sapiens myosin, light polypeptide 6, alkali, smooth muscle and non- muscle (MYL6), transcript variant 1, mRNA NM_04509 Homo sapiens SP110 nuclear body protein (SP110), transcript variant a, mRN. NM_080424 Homo sapiens SP110 nuclear body protein (SP110), transcript variant c, mRN. NM_04510 Homo sapiens SP110 nuclear body protein (SP110), transcript variant b, mRN. NM_04574 Homo sapiens peanut-like 2 (Drosophila) (PNUTL2), transcript variant 1, mRN. NM_080417 Homo sapiens peanut-like 2 (Drosophila) (PNUTL2), transcript variant 3, mRN. NM_080416 Homo sapiens peanut-like 2 (Drosophila) (PNUTL2), transcript variant 2, mRN. NM_080417 Homo sapiens peanut-like 2 (Drosophila) (PNUTL2), transcript variant 2, mRN. NM_080416 Homo sapiens major histocompatibility complex, class I, C (HLA-C), mRNA. NM_080417 Homo sapiens major histocompatibility complex, class I, B (HLA-B), mRNA. NM_080418 Homo sapiens mitochondrion, complete genome NM_080489 Homo sapiens Finkel-Biskis-Reilly murine sarcoma virus (FBR-MuSV)
NM_079837 Homo sapiens BTG3 associated nuclear protein (BANP), transcript variant 2, mRNA NM_017869 Homo sapiens BTG3 associated nuclear protein (BANP), transcript variant 1, mRNA NM_079425 Homo sapiens myosin, light polypeptide 6, alkali, smooth muscle and non-muscle (MYL6), transcript variant 3, mRNA NM_079424 Homo sapiens myosin, light polypeptide 6, alkali, smooth muscle and non-muscle (MYL6), transcript variant 4, mRNA NM_079423 Homo sapiens myosin, light polypeptide 6, alkali, smooth muscle and non-muscle (MYL6), transcript variant 2, mRNA NM_021019 Homo sapiens myosin, light polypeptide 6, alkali, smooth muscle and non-muscle (MYL6), transcript variant 1, mRNA NM_04509 Homo sapiens SP110 nuclear body protein (SP110), transcript variant a, mRN NM_080424 Homo sapiens SP110 nuclear body protein (SP110), transcript variant c, mRN NM_04574 Homo sapiens speanut-like 2 (Drosophila) (PNUTL2), transcript variant 1, mRN NM_080417 Homo sapiens peanut-like 2 (Drosophila) (PNUTL2), transcript variant 3, mRN NM_080416 Homo sapiens peanut-like 2 (Drosophila) (PNUTL2), transcript variant 2, mRN NM_080417 Homo sapiens peanut-like 2 (Drosophila) (PNUTL2), transcript variant 2, mRN NM_080416 Homo sapiens major histocompatibility complex, class I, C (HLA-C), mRNA NM_080417 Homo sapiens major histocompatibility complex, class I, B (HLA-B), mRNA NM_080418 Homo sapiens mitochondrion, complete genome NM_080489 Homo sapiens syndecan binding protein (syntenin) 2 (SDCBP2), mRNA
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NM_079425 Homo sapiens BTG3 associated nuclear protein (BANP), transcript variant 1, mRNA NM_079425 Homo sapiens myosin, light polypeptide 6, alkali, smooth muscle and non-muscle (MYL6), transcript variant 3, mRNA NM_079424 Homo sapiens myosin, light polypeptide 6, alkali, smooth muscle and non-muscle (MYL6), transcript variant 4, mRNA NM_079423 Homo sapiens myosin, light polypeptide 6, alkali, smooth muscle and non-muscle (MYL6), transcript variant 2, mRNA NM_021019 Homo sapiens myosin, light polypeptide 6, alkali, smooth muscle and non-muscle (MYL6), transcript variant 1, mRNA NM_04509 Homo sapiens SP110 nuclear body protein (SP110), transcript variant a, mRN NM_080424 Homo sapiens SP110 nuclear body protein (SP110), transcript variant c, mRN NM_04510 Homo sapiens SP110 nuclear body protein (SP110), transcript variant b, mRN NM_04574 Homo sapiens peanut-like 2 (Drosophila) (PNUTL2), transcript variant 1, mRN NM_080417 Homo sapiens peanut-like 2 (Drosophila) (PNUTL2), transcript variant 4, mRN NM_080416 Homo sapiens peanut-like 2 (Drosophila) (PNUTL2), transcript variant 3, mRN NM_080415 Homo sapiens major histocompatibility complex, class I, C (HLA-C), mRNA NM_005514 Homo sapiens major histocompatibility complex, class I, B (HLA-B), mRNA NC_001807 Homo sapiens mitochondrion, complete genome NM_080489 Homo sapiens syndecan binding protein (syntenin) 2 (SDCBP2), mRNA
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NM_079423 Homo sapiens myosin, light polypeptide 6, alkali, smooth muscle and non-muscle (MYL6), transcript variant 2, mRNA NM_021019 Homo sapiens myosin, light polypeptide 6, alkali, smooth muscle and non-muscle (MYL6), transcript variant 1, mRNA NM_004509 Homo sapiens SP110 nuclear body protein (SP110), transcript variant a, mRN. NM_080424 Homo sapiens SP110 nuclear body protein (SP110), transcript variant c, mRN. NM_004510 Homo sapiens SP110 nuclear body protein (SP110), transcript variant b, mRN. NM_004574 Homo sapiens peanut-like 2 (Drosophila) (PNUTL2), transcript variant 1, mRN. NM_080417 Homo sapiens peanut-like 2 (Drosophila) (PNUTL2), transcript variant 4, mRN. NM_080416 Homo sapiens peanut-like 2 (Drosophila) (PNUTL2), transcript variant 3, mRN. NM_080415 Homo sapiens major histocompatibility complex, class I, C (HLA-C), mRNA. NM_005514 Homo sapiens major histocompatibility complex, class I, B (HLA-B), mRNA. NC_001807 Homo sapiens mitochondrion, complete genome NM_080489 Homo sapiens syndecan binding protein (syntenin) 2 (SDCBP2), mRNA.
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NM_021019 Homo sapiens myosin, light polypeptide 6, alkali, smooth muscle and non-muscle (MYL6), transcript variant 1, mRNA NM_004509 Homo sapiens SP110 nuclear body protein (SP110), transcript variant a, mRN. NM_080424 Homo sapiens SP110 nuclear body protein (SP110), transcript variant c, mRN. NM_004510 Homo sapiens SP110 nuclear body protein (SP110), transcript variant b, mRN. NM_004574 Homo sapiens peanut-like 2 (Drosophila) (PNUTL2), transcript variant 1, mRN. NM_080417 Homo sapiens peanut-like 2 (Drosophila) (PNUTL2), transcript variant 4, mRN. NM_080416 Homo sapiens peanut-like 2 (Drosophila) (PNUTL2), transcript variant 3, mRN. NM_080415 Homo sapiens peanut-like 2 (Drosophila) (PNUTL2), transcript variant 2, mRN. NM_002117 Homo sapiens major histocompatibility complex, class I, C (HLA-C), mRNA. NM_005514 Homo sapiens major histocompatibility complex, class I, B (HLA-B), mRNA. NC_001807 Homo sapiens mitochondrion, complete genome NM_080489 Homo sapiens syndecan binding protein (syntenin) 2 (SDCBP2), mRNA.
muscle (MYL6), transcript variant 1, mRNA NM 004509 Homo sapiens SP110 nuclear body protein (SP110), transcript variant a, mRN. NM 080424 Homo sapiens SP110 nuclear body protein (SP110), transcript variant c, mRN. NM 004510 Homo sapiens SP110 nuclear body protein (SP110), transcript variant b, mRN. NM 004574 Homo sapiens peanut-like 2 (Drosophila) (PNUTL2), transcript variant 1, mRN. NM 080417 Homo sapiens peanut-like 2 (Drosophila) (PNUTL2), transcript variant 4, mRN. NM 080416 Homo sapiens peanut-like 2 (Drosophila) (PNUTL2), transcript variant 3, mRN. NM 080415 Homo sapiens peanut-like 2 (Drosophila) (PNUTL2), transcript variant 2, mRN. NM 002117 Homo sapiens major histocompatibility complex, class I, C (HLA-C), mRNA. NM 005514 Homo sapiens major histocompatibility complex, class I, B (HLA-B), mRNA. NC 001807 Homo sapiens mitochondrion, complete genome NM 080489 Homo sapiens syndecan binding protein (syntenin) 2 (SDCBP2), mRNA.
NM_004509 Homo sapiens SP110 nuclear body protein (SP110), transcript variant a, mRN. NM_080424 Homo sapiens SP110 nuclear body protein (SP110), transcript variant c, mRN. NM_004510 Homo sapiens SP110 nuclear body protein (SP110), transcript variant b, mRN. NM_004574 Homo sapiens peanut-like 2 (Drosophila) (PNUTL2), transcript variant 1, mRN. NM_080417 Homo sapiens peanut-like 2 (Drosophila) (PNUTL2), transcript variant 4, mRN. NM_080416 Homo sapiens peanut-like 2 (Drosophila) (PNUTL2), transcript variant 3, mRN. NM_080415 Homo sapiens peanut-like 2 (Drosophila) (PNUTL2), transcript variant 2, mRN. NM_080415 Homo sapiens major histocompatibility complex, class I, C (HLA-C), mRNA. NM_005514 Homo sapiens major histocompatibility complex, class I, B (HLA-B), mRNA. NC_001807 Homo sapiens mitochondrion, complete genome NM_080489 Homo sapiens syndecan binding protein (syntenin) 2 (SDCBP2), mRNA.
NM_080424 Homo sapiens SP110 nuclear body protein (SP110), transcript variant c, mRN. NM_004510 Homo sapiens SP110 nuclear body protein (SP110), transcript variant b, mRN. NM_004574 Homo sapiens peanut-like 2 (Drosophila) (PNUTL2), transcript variant 1, mRN. NM_080417 Homo sapiens peanut-like 2 (Drosophila) (PNUTL2), transcript variant 4, mRN. NM_080416 Homo sapiens peanut-like 2 (Drosophila) (PNUTL2), transcript variant 3, mRN. NM_080415 Homo sapiens peanut-like 2 (Drosophila) (PNUTL2), transcript variant 2, mRN. NM_080415 Homo sapiens major histocompatibility complex, class I, C (HLA-C), mRNA. NM_005514 Homo sapiens major histocompatibility complex, class I, B (HLA-B), mRNA. NC_001807 Homo sapiens mitochondrion, complete genome NM_080489 Homo sapiens syndecan binding protein (syntenin) 2 (SDCBP2), mRNA.
NM 004510 Homo sapiens SP110 nuclear body protein (SP110), transcript variant b, mRN NM 004574 Homo sapiens peanut-like 2 (Drosophila) (PNUTL2), transcript variant 1, mR1 NM 080417 Homo sapiens peanut-like 2 (Drosophila) (PNUTL2), transcript variant 4, mR1 NM 080416 Homo sapiens peanut-like 2 (Drosophila) (PNUTL2), transcript variant 3, mR1 NM 080415 Homo sapiens peanut-like 2 (Drosophila) (PNUTL2), transcript variant 2, mR1 NM 002117 Homo sapiens major histocompatibility complex, class I, C (HLA-C), mRNA NM 005514 Homo sapiens major histocompatibility complex, class I, B (HLA-B), mRNA NC 001807 Homo sapiens mitochondrion, complete genome NM 080489 Homo sapiens syndecan binding protein (syntenin) 2 (SDCBP2), mRNA
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NC 001807 Homo sapiens mitochondrion, complete genome NM 080489 Homo sapiens syndecan binding protein (syntenin) 2 (SDCBP2), mRNA
NM_080489 Homo sapiens syndecan binding protein (syntenin) 2 (SDCBP2), mRNA
NM_001997 Homo sapiens Finkel-Biskis-Reilly murine sarcoma virus (FBR-MuSV)
ubiquitously expressed (fox derived); ribosomal protein S30 (FAU), mRNA
NM_057179 Homo sapiens likely ortholog of mouse and rat twist-related bHLH protein
Dermo-1 (DERMO1), mRNA
NM_001008 Homo sapiens ribosomal protein S4, Y-linked (RPS4Y), mRNA
NM_001007 Homo sapiens ribosomal protein S4, X-linked (RPS4X), mRNA NM_005192 Homo sapiens cyclin-dependent kinase inhibitor 3 (CDK2-associated dual
NM_005192 Homo sapiens cyclin-dependent kinase inhibitor 3 (CDK2-associated dual specificity phosphatase) (CDKN3), mRNA
NM_079421 Homo sapiens cyclin-dependent kinase inhibitor 2D (p19, inhibits CDK4)
(CDKN2D), transcript variant 2, mRNA
NM_001800 Homo sapiens cyclin-dependent kinase inhibitor 2D (p19, inhibits CDK4)
(CDKN2D), transcript variant 1, mRNA
NM_078626 Homo sapiens cyclin-dependent kinase inhibitor 2C (p18, inhibits CDK4)
(CDKN2C), transcript variant 2, mRNA
NM_001262 Homo sapiens cyclin-dependent kinase inhibitor 2C (p18, inhibits CDK4)
(CDKN2C), transcript variant 1, mRNA
NM_078487 Homo sapiens cyclin-dependent kinase inhibitor 2B (p15, inhibits CDK4)
(CDKN2B), transcript variant 2, mRNA
NM_004936 Homo sapiens cyclin-dependent kinase inhibitor 2B (p15, inhibits CDK4)
(CDKN2B), transcript variant 1, mRNA
NM_004896 Homo sapiens vacuolar protein sorting 26 (yeast) (VPS26), mRNA
NM_052945 Homo sapiens BAFF receptor (BAFFR), mRNA
NM_022648 Homo sapiens tensin (TNS), mRNA
NM_078480 Homo sapiens fuse-binding protein-interacting repressor (SIAHBP1), transcrip
variant 1, mRNA
NM_014281 Homo sapiens fuse-binding protein-interacting repressor (SIAHBP1), transcrip
variant 2, mRNA

2, mRNA NM_078471 Homo sapiens TGFB1-induced anti-apoptotic factor 1 (TIAF1), transcript variant 1, mRNA NM_001852 Homo sapiens collagen, type IX, alpha 2 (COL9A2), mRNA NM_078485 Homo sapiens collagen, type IX, alpha 1 (COL9A1), transcript variant 2, mRNA NM_001851 Homo sapiens collagen, type IX, alpha 1 (COL9A1), transcript variant 1, mRNA NM_001851 Homo sapiens collagen, type IX, alpha 1 (COL9A1), transcript variant 1, mRNA NM_01354 Homo sapiens CCR4-NOT transcription complex, subunit 7 (CNOT7), transcript variant 1, mRNA NM_001384 Homo sapiens CCR4-NOT transcription complex, subunit 7 (CNOT7), transcript variant 1, mRNA NM_004064 Homo sapiens cyclin-dependent kinase inhibitor 1B (p27, Kip1) (CDKN1B), mRNA NM_00389 Homo sapiens cyclin-dependent kinase inhibitor 1A (p21, Cip1) (CDKN1A), transcript variant 1, mRNA NM_078467 Homo sapiens cyclin-dependent kinase inhibitor 1A (p21, Cip1) (CDKN1A), transcript variant 2, mRNA NM_003936 Homo sapiens cDK2-associated protein 1 (CDK2AP1), mRNA NM_078481 Homo sapiens CDK2-associated protein 1 (CDK2AP1), mRNA NM_078481 Homo sapiens CD97 antigen (CD97), transcript variant 1, mRNA NM_080432 Homo sapiens vacuolar protein sorting protein 18 (VPS18), transcript variant 2, mRNA NM_080432 Homo sapiens vacuolar protein sorting protein 18 (VPS18), transcript variant 1, mRNA NM_080431 Homo sapiens vacuolar protein sorting 16 (yeast) (VPS16), transcript variant 1, mRNA NM_080414 Homo sapiens vacuolar protein sorting 16 (yeast) (VPS16), transcript variant 1, mRNA NM_080415 Homo sapiens vacuolar protein sorting 11 (yeast) (VPS16), transcript variant 1, mRNA NM_080410 Homo sapiens sequin 12 (CCN72), transcript variant 1, mRNA NM_080410 Homo sapiens sequin 12 (CCN72), transcript variant 1, mRNA NM_080411 Homo sapiens sequin 12 (CCN72), transcript variant 1, mRNA NM_080412 Homo sapiens sequin 12 (CCN72), transcript variant 1, mRNA NM_080404 Homo sapiens sequin 12 (CCN72), transcript variant 1, mRNA NM_080404 Homo sapiens cyclin 12 (CCN712), transcript variant 1, mRNA NM_08047 Homo sapie		
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NM_012106 Homo sapiens binder of Arl Two (BART1), mRNA NM_006095 Homo sapiens ATPase, aminophospholipid transporter (APLT), Class I, type 8A, member 1 (ATP8A1), mRNA NM_058241 Homo sapiens cyclin T2 (CCNT2), transcript variant b, mRNA NM_001241 Homo sapiens cyclin T2 (CCNT2), transcript variant a, mRNA NM_001240 Homo sapiens cyclin T1 (CCNT1), mRNA NM_000474 Homo sapiens twist homolog (acrocephalosyndactyly 3; Saethre-Chotzen syndrome) (Drosophila) (TWIST), mRNA NM_080475 Homo sapiens serine (or cysteine) proteinase inhibitor, clade B (ovalbumin), member 11 (SERPINB11), mRNA NM_021209 Homo sapiens caspase recruitment domain protein 12 (CARD12), mRNA NM_014550 Homo sapiens caspase recruitment domain protein 10 (CARD10), mRNA NM_012287 Homo sapiens centaurin, beta 2 (CENTB2), mRNA NM_007049 Homo sapiens butyrophilin, subfamily 2, member A1 (BTN2A1), transcript variant 1, mRNA NM_078476 Homo sapiens butyrophilin, subfamily 2, member A1 (BTN2A1), transcript variant 2, mRNA	NM_005806	
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member 11 (SERPINB11), mRNA NM_021209 Homo sapiens caspase recruitment domain protein 12 (CARD12), mRNA NM_014550 Homo sapiens caspase recruitment domain protein 10 (CARD10), mRNA NM_012287 Homo sapiens centaurin, beta 2 (CENTB2), mRNA NM_007049 Homo sapiens butyrophilin, subfamily 2, member A1 (BTN2A1), transcript variant 1, mRNA NM_078476 Homo sapiens butyrophilin, subfamily 2, member A1 (BTN2A1), transcript variant 2, mRNA	NM 080475	Homo saniens serine (or systeine) proteiness inhibitor slade B (overthermin)
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NM_078476 Homo sapiens butyrophilin, subfamily 2, member A1 (BTN2A1), transcript variant 2, mRNA	NM_007049	
	NM_078476	Homo sapiens butyrophilin, subfamily 2, member A1 (BTN2A1), transcript
	NM_004444	Homo sapiens EphB4 (EPHB4), mRNA

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NM_004443	Homo sapiens EphB3 (EPHB3), mRNA		
NM_004442	Homo sapiens EphB2 (EPHB2), transcript variant 1, mRNA		
NM_017449	Homo sapiens EphB2 (EPHB2), transcript variant 2, mRNA		
NM_004535	Homo sapiens my lin transcription factor 1 (MYT1), mRNA		
NM_006800	Homo sapiens male-specific lethal 3-like 1 (Drosophila) (MSL3L1), transcript variant 3, mRNA		
NM_078630	Homo sapiens male-specific lethal 3-like 1 (Drosophila) (MSL3L1), transcript variant 2, mRNA		
NM_078629	Homo sapiens male-specific lethal 3-like 1 (Drosophila) (MSL3L1), transcript variant 1, mRNA		
NM_078628	Homo sapiens male-specific lethal 3-like 1 (Drosophila) (MSL3L1), transcript variant 4, mRNA		
NM_080431	Homo sapiens actin related protein M2 (ARPM2), mRNA		
NM_080430	Homo sapiens selenoprotein SelM (SELM), mRNA		
NM_052944	Homo sapiens putative sodium-coupled cotransporter RKST1 (RKST1), mRNA		
NM_024831	Homo sapiens nuclear receptor coactivator 6 interacting protein (NCOA6IP), mRNA		
NM_032803	Homo sapiens solute carrier family 7 (cationic amino acid transporter, y+ system), member 3 (SLC7A3), mRNA		
NM_080385	Homo sapiens carboxypeptidase A5 (CPA5), mRNA		
NM_016476	Homo sapiens APC11 anaphase promoting complex subunit 11 homolog (yeast) (ANAPC11), mRNA		
NM_080389	Homo sapiens defensin, beta 4 (DEFB4), mRNA		
NM_032646	Homo sapiens tweety homolog 2 (Drosophila) (TTYH2), mRNA		
NM 006928	Homo sapiens silver homolog (mouse) (SILV), mRNA		
NM 080390	Homo sapiens my048 protein (my048), mRNA		
NM 080388	Homo sapiens hypothetical protein MGC17528 (MGC17528), mRNA		
NM 080387	Homo sapiens C-type lectin-like receptor (CLEC-6), mRNA		
NM_080284	Homo sapiens ATP-binding cassette, sub-family A (ABC1), member 6 (ABCA6), mRNA		
NM_080283	Homo sapiens ATP-binding cassette, sub-family A (ABC1), member 9 (ABCA9), mRNA		
NM_080282	Homo sapiens ATP-binding cassette, sub-family A (ABC1), member 10 (ABCA10), mRNA		
NM_006549	Homo sapiens calcium/calmodulin-dependent protein kinase kinase 2, beta (CAMKK2), mRNA		
NM_007200	Homo sapiens A kinase (PRKA) anchor protein 13 (AKAP13), mRNA		
NM_002476	Homo sapiens myosin, light polypeptide 4, alkali; atrial, embryonic (MYL4), mRNA		
NM_001853	Homo sapiens collagen, type IX, alpha 3 (COL9A3), mRNA		
NM_006001	Homo sapiens tubulin, alpha 2 (TUBA2), transcript variant 1, mRNA		
NM_079836	Homo sapiens tubulin, alpha 2 (TUBA2), transcript variant 2, mRNA		
NM_006000	Homo sapiens tubulin, alpha 1 (testis specific) (TUBA1), mRNA		
NM_004376	Homo sapiens COX15 homolog, cytochrome c oxidase assembly protein (yeast) (COX15), nuclear gene encoding mitochondrial protein, transcript variant 2, mRNA		
NM_024407	Homo sapiens NADH dehydrogenase (ubiquinone) Fe-S protein 7 (20kD) (NADH-coenzyme Q reductase) (NDUFS7), mRNA		
NM 078625	Homo sapiens vanin 3 (VNN3), transcript variant 2, mRNA		
NM 018399	Homo sapiens vanin 3 (VNN3), transcript variant 1, mRNA		
NM 078488	Homo sapiens vanin 2 (VNN2), transcript variant 1, mRNA Homo sapiens vanin 2 (VNN2), transcript variant 2, mRNA		
NM 004665	Homo sapiens vanin 2 (VNN2), transcript variant 1, mRNA		
1111 00 1000	Tromo sapiens vann 2 (viviv2), nanscript variant 1, mkivA		

NM_013245	Homo sapiens vacuolar protein sorting factor 4A (VPS4A), mRNA
NM_058240	Homo sapiens solute carrier family 8 (sodium-calcium exchanger), member 3
	(SLC8A3), transcript variant b, mRNA
NM_033262	Homo sapiens solute carrier family 8 (sodium-calcium exchanger), member 3
	(SLC8A3), transcript variant a, mRNA
NM_004869	Homo sapiens suppressor of K+ transport defect 1 (SKD1), mRNA
NM_078474	Homo sapiens BBP-like protein 2 (BLP2), transcript variant 1, mRNA
NM_025141	Homo sapiens BBP-like protein 2 (BLP2), transcript variant 2, mRNA
NM_078473	Homo sapiens BBP-like protein 1 (BLP1), transcript variant 1, mRNA
NM_031940	Homo sapiens BBP-like protein 1 (BLP1), transcript variant 2, mRNA
NM_020749	Homo sapiens AT2 receptor-interacting protein 1 (ATIP1), mRNA
NM_018672	Homo sapiens ATP-binding cassette, sub-family A (ABC1), member 5
	(ABCA5), mRNA
NM_020177	Homo sapiens feminization 1 homolog a (FEM1A), mRNA
NM_002088	Homo sapiens glutamate receptor, ionotropic, kainate 5 (GRIK5), mRNA
NM_006835	Homo sapiens cyclin I (CCNI), mRNA
NM_001239	Homo sapiens cyclin H (CCNH), mRNA
NM_014286	Homo sapiens frequenin homolog (Drosophila) (FREQ), mRNA
NM_006650	Homo sapiens complexin 2 (CPLX2), mRNA
NM_006651	Homo sapiens complexin 1 (CPLX1), mRNA
NM_006463	Homo sapiens associated molecule with the SH3 domain of STAM (AMSH),
	mRNA
NM_001850	Homo sapiens collagen, type VIII, alpha 1 (COL8A1), mRNA
NM_000094	Homo sapiens collagen, type VII, alpha 1 (epidermolysis bullosa, dystrophic,
	dominant and recessive) (COL7A1), mRNA
NM_000077	Homo sapiens cyclin-dependent kinase inhibitor 2A (melanoma, p16, inhibits
	CDK4) (CDKN2A), transcript variant 1, mRNA
NM_058197	Homo sapiens cyclin-dependent kinase inhibitor 2A (melanoma, p16, inhibits
20.0000	CDK4) (CDKN2A), transcript variant 3, mRNA
NM_058196	Homo sapiens cyclin-dependent kinase inhibitor 2A (melanoma, p16, inhibits
27.650105	CDK4) (CDKN2A), transcript variant 2, mRNA
NM_058195	Homo sapiens cyclin-dependent kinase inhibitor 2A (melanoma, p16, inhibits
NB4 014000	CDK4) (CDKN2A), transcript variant 4, mRNA
NM_014800	Homo sapiens engulfment and cell motility 1 (ced-12 homolog, C. elegans)
NM 079834	(ELMO1), mRNA
NM 019110	Homo sapiens secretory carrier membrane protein 4 (SCAMP-4), mRNA
	Homo sapiens hypothetical protein P1 p373c6 (P1P373C6), mRNA
NM_022086	Homo sapiens engulfment and cell motility 2 (ced-12 homolog, C. elegans) (ELMO2), mRNA
NM 058183	Homo sapiens SON DNA binding protein (SON), mRNA
NM_003103	Homo sapiens SON DNA binding protein (SON), mRNA Homo sapiens SON DNA binding protein (SON), mRNA
NM_030767	Homo sapiens AT-hook transcription factor AKNA (AKNA), mRNA
NM_058191	Homo sapiens A1-nook transcription factor AKNA (AKNA), mRNA Homo sapiens chromosome 21 open reading frame 66 (C21orf66), mRNA
NM 015657	Homo sapiens ATP-binding cassette, sub-family A (ABC1), member 12
1441_012027	(ABCA12), mRNA
NM 020427	Homo sapiens ARS component B (ARS), mRNA
NM_021638	Homo sapiens actin filament associated protein (AFAP), mRNA
NM_005782	Homo sapiens transcriptional coactivator (ALY), mRNA
NM 031916	Homo sapiens AKAP-associated sperm protein (ASP), mRNA
NM 024083	Homo sapiens alveolar soft part sarcoma chromosome region, candidate 1
	(ASPSCR1), mRNA
NM_058230	Homo sapiens zinc finger protein 354B (ZNF354B), mRNA

NM_021935	Homo sapiens homolog of mouse Bv8 (Bombina variegata 8 kDa); prokineticin 2 precursor (BV8), mRNA
NM 015399	Homo sapiens breast cancer metastasis-suppressor 1 (BRMS1), mRNA
NM 007073	Homo sapiens blood vessel epicardial substance (BVES), mRNA
NM_017726	Homo sapiens protein phosphatase 1, regulatory (inhibitor) subunit 14D (PPP1R14D), mRNA
NM_006451	Homo sapiens polyadenylate binding protein-interacting protein 1 (PAIP1), mRNA
NM_018073	Homo sapiens SSA protein SS-56 (SS-56), mRNA
NM 032812	Homo sapiens tumor endothelial marker 7-related precursor (TEM7R), mRNA
NM_022748	Homo sapiens tumor endothelial marker 6 (TEM6), mRNA
NM_032777	Homo sapiens tumor endothelial marker 5 precursor (TEM5), mRNA
NM_022779	Homo sapiens DEAD/H (Asp-Glu-Ala-Asp/His) box polypeptide 31 (DDX31), mRNA
NM_018454	Homo sapiens nucleolar protein ANKT (ANKT), mRNA
NM_016489	Homo sapiens uridine 5' monophosphate hydrolase 1 (UMPH1), mRNA
NM_078483	Homo sapiens lysosomal amino acid transporter 1 (LYAAT1), mRNA
NM_019606	Homo sapiens hypothetical protein FLJ20257 (FLJ20257), mRNA
NM_015256	Homo sapiens fatty-acid-Coenzyme A ligase, long-chain 6 (FACL6), mRNA
NM_003393	Homo sapiens wingless-type MMTV integration site family, member 8B (WNT8B), mRNA
NM_058244	Homo sapiens wingless-type MMTV integration site family, member 8A (WNT8A), transcript variant 2, mRNA
NM_058238	Homo sapiens wingless-type MMTV integration site family, member 7B (WNT7B), mRNA
NM_004625	Homo sapiens wingless-type MMTV integration site family, member 7A (WNT7A), mRNA
NM 058242	Homo sapiens keratin 6C (KRT6C), mRNA
NM 005555	Homo sapiens keratin 6B (KRT6B), mRNA
NM_005554	Homo sapiens keratin 6A (KRT6A), mRNA
NM_058207	Homo sapiens sperm associated antigen 11 (SPAG11), transcript variant E, mRNA
NM_058206	Homo sapiens sperm associated antigen 11 (SPAG11), transcript variant B, mRNA
NM_058203	Homo sapiens sperm associated antigen 11 (SPAG11), transcript variant C, mRNA
NM_058202	Homo sapiens sperm associated antigen 11 (SPAG11), transcript variant H, mRNA
NM_058201	Homo sapiens sperm associated antigen 11 (SPAG11), transcript variant D, mRNA
NM_058200	Homo sapiens sperm associated antigen 11 (SPAG11), transcript variant G, mRNA
NM_016512	Homo sapiens sperm associated antigen 11 (SPAG11), transcript variant A, mRNA
NM_057180	Homo sapiens vacuolar protein sorting 29 (yeast) (VPS29), transcript variant 2, mRNA
NM_016226	Homo sapiens vacuolar protein sorting 29 (yeast) (VPS29), transcript variant 1, mRNA
NM_053004	Homo sapiens guanine nucleotide binding protein (G protein), beta polypeptide 1-like (GNB1L), mRNA
NM_003902	Homo sapiens far upstream element (FUSE) binding protein 1 (FUBP1), mRNA
NM_058217	Homo sapiens RAD51 homolog C (S. cerevisiae) (RAD51C), transcript variant

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NM_058216	Homo sapiens RAD51 homolog C (S. cerevisiae) (RAD51C), transcript variant 1, mRNA
NM_002876	Homo sapiens RAD51 homolog C (S. cerevisiae) (RAD51C), transcript variant 2, mRNA
NM_058179	Homo sapiens phosphoserine aminotransferase (PSA), transcript variant 1, mRNA
NM_021154	Homo sapiens phosphoserine aminotransferase (PSA), transcript variant 2, mRNA
NM_078469	Homo sapiens BRCA2 and CDKN1A interacting protein (BCCIP), transcript variant C, mRNA
NM_078468	Homo sapiens BRCA2 and CDKN1A interacting protein (BCCIP), transcript variant B, mRNA
NM_016567	Homo sapiens BRCA2 and CDKN1A interacting protein (BCCIP), transcript variant A, mRNA
NM_058177	Homo sapiens histone deacetylase 9 (HDAC9-PENDING), transcript variant 2, mRNA
NM_058176	Homo sapiens histone deacetylase 9 (HDAC9-PENDING), transcript variant 1, mRNA
NM 022110	Homo sapiens FK506 binding protein like (FKBPL), mRNA
NM 012181	Homo sapiens FK506 binding protein 8 (38kD) (FKBP8), mRNA
NM 003602	Homo sapiens FK506 binding protein 6 (36kD) (FKBP6), mRNA
NM 004117	Homo sapiens FK506 binding protein 5 (FKBP5), mRNA
NM 002014	Homo sapiens FK506 binding protein 4 (59kD) (FKBP4), mRNA
NM_057092	Homo sapiens FK506 binding protein 2 (13kD) (FKBP2), transcript variant 2, mRNA
NM_004470	Homo sapiens FK506 binding protein 2 (13kD) (FKBP2), transcript variant 1, mRNA
NM_004116	Homo sapiens FK506 binding protein 1B (12.6 kD) (FKBP1B), transcript variant 1, mRNA
NM_054033	Homo sapiens FK506 binding protein 1B (12.6 kD) (FKBP1B), transcript variant 2, mRNA
NM_000801	Homo sapiens FK506 binding protein 1A (12kD) (FKBP1A), transcript variant 12B, mRNA
NM_054014	Homo sapiens FK506 binding protein 1A (12kD) (FKBP1A), transcript variant 12A, mRNA
NM_057175	Homo sapiens hypothetical protein FLJ13340 (FLJ13340), transcript variant 1, mRNA
NM_025085	Homo sapiens hypothetical protein FLJ13340 (FLJ13340), transcript variant 2, mRNA
NM_014708	Homo sapiens kinetochore associated 1 (KNTC1), mRNA
NM_058199	Homo sapiens olfactomedin 1 (OLFM1), transcript variant 3, mRNA
NM_014279	Homo sapiens olfactomedin 1 (OLFM1), transcript variant 1, mRNA
NM_057174	Homo sapiens peroxisomal biogenesis factor 16 (PEX16), transcript variant 2, mRNA
NM_033118	Homo sapiens myosin light chain kinase 2, skeletal muscle (MYLK2), mRNA
NM_019117	Homo sapiens kelch-like 4 (Drosophila) (KLHL4), transcript variant 1, mRNA
NM_005103	Homo sapiens fasciculation and elongation protein zeta 1 (zygin I) (FEZ1), transcript variant 1, mRNA
NM_022549	Homo sapiens fasciculation and elongation protein zeta 1 (zygin I) (FEZ1), transcript variant 2, mRNA
NM_005112	Homo sapiens WD repeat domain 1 (WDR1), transcript variant 2, mRNA

Homo sapiens WD repeat domain 1 (WDR1), transcript variant 1, mRNA
Homo sapiens cytochrome c oxidase subunit Vb (COX5B), nuclear gene
encoding mitochondrial protein, mRNA
Homo sapiens cytochrome c oxidase subunit Va (COX5A), nuclear gene
encoding mitochondrial protein, mRNA
Homo sapiens kelch-like 4 (Drosophila) (KLHL4), transcript variant 2, mRNA
Homo sapiens cortactin binding protein 2 (CORTBP2), mRNA
Homo sapiens cyclin-dependent kinase 7 (MO15 homolog, Xenopus laevis, cdk-activating kinase) (CDK7), mRNA
Homo sapiens adaptor-related protein complex 1, sigma 1 subunit (AP1S1), transcript variant 2, mRNA
Homo sapiens adaptor-related protein complex 1, sigma 1 subunit (AP1S1), transcript variant 1, mRNA
Homo sapiens unc-119 homolog (C. elegans) (UNC119), transcript variant 1, mRNA
Homo sapiens unc-119 homolog (C. elegans) (UNC119), transcript variant 2, mRNA
Homo sapiens protocadherin LKC (PC-LKC), mRNA
Homo sapiens mitogen-activated protein kinase kinase kinase 3 (MAP3K3), mRNA
Homo sapiens unc-5 homolog B (C. elegans) (UNC5C), mRNA
Homo sapiens angiopoietin-like 1 (ANGPTL1), mRNA
Homo sapiens FUS interacting protein (serine-arginine rich) 1 (FUSIP1), transcript variant 2, mRNA
Homo sapiens FUS interacting protein (serine-arginine rich) 1 (FUSIP1), transcript variant 1, mRNA
Homo sapiens ankylosis, progressive homolog (mouse) (ANKH), transcript variant 2, mRNA
Homo sapiens ankylosis, progressive homolog (mouse) (ANKH), transcript variant 1, mRNA
Homo sapiens Sec23 homolog B (S. cerevisiae) (SEC23B), transcript variant 1, mRNA
Homo sapiens Sec23 homolog B (S. cerevisiae) (SEC23B), transcript variant 3, mRNA
Homo sapiens Sec23 homolog B (S. cerevisiae) (SEC23B), transcript variant 2, mRNA
Homo sapiens tektin 1 (TEKT1), mRNA
Homo sapiens phosphoprotein associated with glycosphingolipid-enriched microdomains (PAG), mRNA
Homo sapiens ADAM-like, decysin 1 (ADAMDEC1), mRNA
Homo sapiens immediate early response 5 (IER5), mRNA
Homo sapiens coronin, actin binding protein, 2A (CORO2A), transcript variant 2, mRNA
Homo sapiens coronin, actin binding protein, 2A (CORO2A), transcript variant 1, mRNA
Homo sapiens caspase recruitment domain family, member 6 (CARD6), mRNA
Homo sapiens caspase recruitment domain family, member 9 (CARD9), transcript variant 2, mRNA
Homo sapiens caspase recruitment domain family, member 9 (CARD9), transcript variant 1, mRNA
Homo sapiens caspase recruitment domain family, member 9 (CARD9), transcript variant 3, mRNA

NM_052978	Homo sapiens tripartite motif-containing 9 (TRIM9), transcript variant 2, mRNA
NM_015163	Homo sapiens tripartite motif-containing 9 (TRIM9), transcript variant 1, mRNA
NM_052840	Homo sapiens bruno-like 6, RNA binding protein (Drosophila) (BRUNOL6), mRNA
NM_000967	Homo sapiens ribosomal protein L3 (RPL3), mRNA
NM_015125	Homo sapiens capicua homolog (Drosophila) (CIC), mRNA
NM_018256	Homo sapiens WD repeat domain 12 (WDR12), mRNA
NM_016601	Homo sapiens potassium channel, subfamily K, member 9 (TASK-3) (KCNK9), mRNA
NM_033415	Homo sapiens hypothetical gene MGC19595 (MGC19595), mRNA
NM_001253	Homo sapiens CDC5 cell division cycle 5-like (S. pombe) (CDC5L), mRNA
NM_007065	Homo sapiens CDC37 cell division cycle 37 homolog (S. cerevisiae) (CDC37), mRNA
NM_003504	Homo sapiens CDC45 cell division cycle 45-like (S. cerevisiae) (CDC45L), mRNA
NM_006035	Homo sapiens CDC42 binding protein kinase beta (DMPK-like) (CDC42BPB), mRNA
NM_044472	Homo sapiens cell division cycle 42 (GTP binding protein, 25kD) (CDC42), transcript variant 2, mRNA
NM_001791	Homo sapiens cell division cycle 42 (GTP binding protein, 25kD) (CDC42), transcript variant 1, mRNA
NM_001254	Homo sapiens CDC6 cell division cycle 6 homolog (S. cerevisiae) (CDC6), mRNA
NM_022894	Homo sapiens poly(A) polymerase gamma (PAPOLG), mRNA
NM_033655	Homo sapiens cell recognition molecule CASPR3 (CASPR3), transcript variant 1, mRNA
NM_024879	Homo sapiens cell recognition molecule CASPR3 (CASPR3), transcript variant 2, mRNA
NM 012115	Homo sapiens CASP8 associated protein 2 (CASP8AP2), mRNA
NM 012173	Homo sapiens F-box only protein 25 (FBXO25), mRNA
NM_033624	Homo sapiens F-box only protein 21 (FBXO21), transcript variant 1, mRNA
NM_015002	Homo sapiens F-box only protein 21 (FBXO21), transcript variant 2, mRNA
NM 033625	Homo sapiens ribosomal protein L34 (RPL34), transcript variant 2, mRNA
NM 000995	Homo sapiens ribosomal protein L34 (RPL34), transcript variant 1, mRNA
NM_033540	Homo sapiens mitofusin 1 (MFN1), transcript variant 1, mRNA
NM 005612	Homo sapiens RE1-silencing transcription factor (REST), mRNA
NM 007085	Homo sapiens follistatin-like 1 (FSTL1), mRNA
NM 000993	Homo sapiens ribosomal protein L31 (RPL31), mRNA
NM 012180	Homo sapiens F-box only protein 8 (FBXO8), mRNA
NM 033182	Homo sapiens F-box protein FBX30 (FBX30), mRNA
NM_033406	Homo sapiens F-box only protein 3 (FBXO3), transcript variant 2, mRNA
NM 012175	Homo sapiens F-box only protein 3 (FBXO3), transcript variant 1, mRNA
NM_017425	Homo sapiens sperm autoantigenic protein 17 (SPA17), mRNA
NM_005633	Homo sapiens son of sevenless homolog 1 (Drosophila) (SOS1), mRNA
NM_003333	Homo sapiens ubiquitin A-52 residue ribosomal protein fusion product 1 (UBA52), mRNA
NM 019894	Homo sapiens transmembrane protease, serine 4 (TMPRSS4), mRNA
NM_033313	Homo sapiens CDC14 cell division cycle 14 homolog A (S. cerevisiae) (CDC14A), transcript variant 3, mRNA
NM_033312	Homo sapiens CDC14 cell division cycl 14 homolog A (S. cerevisiae) (CDC14A), transcript variant 2, mRNA
NM_003672	Homo sapiens CDC14 cell division cycle 14 homolog A (S. cerevisiae)

		Les States
		(CDC14A), transcript variant 1, mRNA
	NM_005786	Homo sapiens serologically defined colon cancer antigen 33 (SDCCAG33), mRNA
	NM_003618	Homo sapiens mitogen-activated protein kinase kinase kinase kinase 3 (MAP4K3), mRNA
	NM_006577	Homo sapiens UDP-GlcNAc:betaGal beta-1,3-N-acetylglucosaminyltransferase 1 (B3GNT1), transcript variant 1, mRNA
	NM_020981	Homo sapiens UDP-Gal:betaGlcNAc beta 1,3-galactosyltransferase, polypeptide 1 (B3GALT1), mRNA
	NM_033252	Homo sapiens UDP-GlcNAc:betaGal beta-1,3-N-acetylglucosaminyltransferase 1 (B3GNT1), transcript variant 2, mRNA
	NM 002954	Homo sapiens ribosomal protein S27a (RPS27A), mRNA
	NM 000971	Homo sapiens ribosomal protein L7 (RPL7), mRNA
	NM 033344	Homo sapiens egl nine homolog 3 (C. elegans) (EGLN3), mRNA
	NM 024023	Homo sapiens unkempt-like (Drosophila) (UNKL), mRNA
	NM_033221	Homo sapiens tripartite motif-containing 14 (TRIM14), transcript variant 4, mRNA
	NM_033220	Homo sapiens tripartite motif-containing 14 (TRIM14), transcript variant 3, mRNA
	NM_033219	Homo sapiens tripartite motif-containing 14 (TRIM14), transcript variant 2, mRNA
	NM_014788	Homo sapiens tripartite motif-containing 14 (TRIM14), transcript variant 1, mRNA
	NM_006074	Homo sapiens tripartite motif-containing 22 (TRIM22), mRNA
	NM_012210	Homo sapiens tripartite motif-containing 32 (TRIM32), mRNA
	NM_007276	Homo sapiens chromobox homolog 3 (HP1 gamma homolog, Drosophila) (CBX3), mRNA
	NM 025227	Homo sapiens hypothetical protein dJ726C3.2 (DJ726C3.2), mRNA
• = 4/5π ² 4	NM 015271	Homo sapiens tripartite motif-containing 2 (TRIM2), mRNA
	NM_017838	Homo sapiens nucleolar protein family A, member 2 (H/ACA small nucleolar RNPs) (NOLA2), mRNA
	NM_032993	Homo sapiens nucleolar protein family A, member 1 (H/ACA small nucleolar RNPs) (NOLA1), transcript variant 2, mRNA
	NM_018983	Homo sapiens nucleolar protein family A, member 1 (H/ACA small nucleolar RNPs) (NOLA1), transcript variant 1, mRNA
	NM_004722	Homo sapiens adaptor-related protein complex 4, mu 1 subunit (AP4M1), mRNA
	NM_033066	Homo sapiens membrane protein, palmitoylated 4 (MAGUK p55 subfamily member 4) (MPP4), mRNA
	NM 033030	Homo sapiens bol, boule-like (Drosophila) (BOLL), mRNA
	NM_004216	Homo sapiens death effector domain-containing (DEDD), transcript variant 2, mRNA
	NM_032998	Homo sapiens death effector domain-containing (DEDD), transcript variant 1, mRNA
	NM 033010	Homo sapiens poly(rC) binding protein 4 (PCBP4), transcript variant 4, mRNA
	NM 033009	Homo sapiens poly(rC) binding protein 4 (PCBP4), transcript variant 2, mRNA
	NM 033008	Homo sapiens poly(rC) binding protein 4 (PCBP4), transcript variant 3, mRNA
	NM 020418	Homo sapiens poly(rC) binding protein 4 (PCBP4), transcript variant 1, mRNA
	NM 032944	Homo sapiens serine/threonine kinase 31 (STK31), transcript variant 2, mRNA
	NM 031414	Homo sapiens serine/threonine kinase 31 (STK31), transcript variant 1, mRNA
	NM 014302	Homo sapiens Sec61 gamma (SEC61G), mRNA
	NM_013336	Homo sapiens protein transport protein SEC61 alpha subunit isoform 1

г	(SEC(1A1) mDNA
27.6 001.401	(SEC61A1), mRNA Homo sapiens tethering factor SEC34 (SEC34), mRNA
NM 031431	
NM 015490	Homo sapiens secretory pathway component Sec31B-1 (SEC31B-1), mRNA
NM_004892	Homo sapiens SEC22 vesicle trafficking protein-like 1 (S. cerevisiae)
>7.5 000070	(SEC22L1), mRNA Homo sapiens vesicle trafficking protein (SEC22C), transcript variant 1, mRNA
NM_032970	
NM_000969	Homo sapiens ribosomal protein L5 (RPL5), mRNA
NM_005034	Homo sapiens polymerase (RNA) II (DNA directed) polypeptide K (7.0kD) (POLR2K), mRNA
NM_014459	Homo sapiens protocadherin 17 (PCDH17), mRNA
NM_032961	Homo sapiens protocadherin 10 (PCDH10), transcript variant 1, mRNA
NM 020815	Homo sapiens protocadherin 10 (PCDH10), transcript variant 2, mRNA
NM 031988	Homo sapiens mitogen-activated protein kinase kinase 6 (MAP2K6), transcript
	variant 2, mRNA
NM_002758	Homo sapiens mitogen-activated protein kinase kinase 6 (MAP2K6), transcript variant 1, mRNA
NM 032419	Homo sapiens dom-3 homolog Z (C. elegans) (DOM3Z), transcript variant 1,
_	mRNA
NM_032966	Homo sapiens Burkitt lymphoma receptor 1, GTP binding protein (BLR1), transcript variant 2, mRNA
NM 001716	Homo sapiens Burkitt lymphoma receptor 1, GTP binding protein (BLR1),
_	transcript variant 1, mRNA
NM 004951	Homo sapiens Epstein-Barr virus induced gene 2 (lymphocyte-specific G
_	protein-coupled receptor) (EBI2), mRNA
NM 004874	Homo sapiens BCL2-associated athanogene 4 (BAG4), mRNA
NM_001016	Homo sapiens ribosomal protein S12 (RPS12), mRNA
NM 031994	Homo sapiens ring finger protein 17 (RNF17), transcript variant short, mRNA
NM_031271	Homo sapiens testis expressed sequence 15 (TEX15), mRNA
NM_018995	Homo sapiens Mov1011, Moloney leukemia virus 10-like 1, homolog (mouse) (MOV10L1), mRNA
NM_032510	Homo sapiens par-6 partitioning defective 6 homolog gamma (C. elegans) (PARD6G), mRNA
NM_006704	Homo sapiens suppressor of G2 allele of SKP1, S. cerevisiae, homolog of (SGT1), mRNA
NM_031968	Homo sapiens nuclear prelamin A recognition factor (NARF), transcript variant 2, mRNA
NM_012336	Homo sapiens nuclear prelamin A recognition factor (NARF), transcript variant 1, mRNA
NM 003980	Homo sapiens microtubule-associated protein 7 (MAP7), mRNA
NM 032380	Homo sapiens elongation factor G2 (EFG2), mRNA
NM 032214	Homo sapiens Src-like-adaptor 2 (SLA2), mRNA
NM 020064	Homo sapiens BarH-like 1 (Drosophila) (BARHL1), mRNA
NM 005916	Homo sapiens MCM7 minichromosome maintenance deficient 7 (S. cerevisiae)
	(MCM7), mRNA
NM 004098	Homo sapiens empty spiracles homolog 2 (Drosophila) (EMX2), mRNA
NM 005826	Homo sapiens heterogeneous nuclear ribonucleoprotein R (HNRPR), mRNA
NM_006418	Homo sapiens differentially expressed in hematopoietic lineages (GW112), mRNA
NM 005016	Homo sapiens poly(rC) binding protein 2 (PCBP2), transcript variant 1, mRNA
NM 031989	Homo sapiens poly(rC) binding protein 2 (PCBP2), transcript variant 2, mRNA
NM 006196	Homo sapiens poly(rC) binding protein 1 (PCBP1), mRNA
NM 031844	Homo sapiens heterogeneous nuclear ribonucleoprotein U (scaffold attachment
TATAT 02 1944	Homo saprens neterogeneous modear moducicobrocom o (section attachment

	factor A) (HNRPU), transcript variant 1, mRNA
NM_004501	Homo sapiens heterogeneous nuclear ribonucleoprotein U (scaffold attachment
	factor A) (HNRPU), transcript variant 2, mRNA
NM_004500	Homo sapiens heterogeneous nuclear ribonucleoprotein C (C1/C2) (HNRPC),
	transcript variant 2, mRNA
NM_031314	Homo sapiens heterogeneous nuclear ribonucleoprotein C (C1/C2) (HNRPC),
	transcript variant 1, mRNA
NM_031370	Homo sapiens heterogeneous nuclear ribonucleoprotein D (AU-rich element
	RNA binding protein 1, 37kD) (HNRPD), transcript variant 1, mRNA
NM_031369	Homo sapiens heterogeneous nuclear ribonucleoprotein D (AU-rich element
	RNA binding protein 1, 37kD) (HNRPD), transcript variant 2, mRNA
NM_002138	Homo sapiens heterogeneous nuclear ribonucleoprotein D (AU-rich element
	RNA binding protein 1, 37kD) (HNRPD), transcript variant 3, mRNA
NM_003903	Homo sapiens CDC16 cell division cycle 16 homolog (S. cerevisiae) (CDC16), mRNA
NM_031483	Homo sapiens itchy homolog E3 ubiquitin protein ligase (mouse) (ITCH), mRNA
NM 031907	Homo sapiens ubiquitin specific protease 26 (USP26), mRNA
NM 031866	Homo sapiens frizzled homolog 8 (Drosophila) (FZD8), mRNA
NG 000004	Homo sapiens genomic cytochrome P450, subfamily IIIA (niphedipine oxidase)
	(CYP3A) on chromosome 7
NM_001788	Homo sapiens CDC10 cell division cycle 10 homolog (S. cerevisiae) (CDC10),
_	mRNA
NM_004276	Homo sapiens calcium binding protein 1 (calbrain) (CABP1), transcript variant
	2, mRNA
NM 031205	Homo sapiens calcium binding protein 1 (calbrain) (CABP1), transcript variant
	1, mRNA
NM_000784	Homo sapiens cytochrome P450, subfamily XXVIIA (steroid 27-hydroxylase,
_	cerebrotendinous xanthomatosis), polypeptide 1 (CYP27A1), nuclear gene
	encoding mitochondrial protein, mRNA
NM_031491	Homo sapiens retinol binding protein 5, cellular (RBP5), mRNA
NM_006929	Homo sapiens superkiller viralicidic activity 2-like (S. cerevisiae) (SKIV2L),
_	mRNA
NM_001447	Homo sapiens FAT tumor suppressor homolog 2 (Drosophila) (FAT2), mRNA
NM_007242	Homo sapiens DEAD/H (Asp-Glu-Ala-Asp/His) box polypeptide 19 (DBP5
_	homolog, yeast) (DDX19), mRNA
NM_006773	Homo sapiens DEAD/H (Asp-Glu-Ala-Asp/His) box polypeptide 18 (Myc-
_	regulated) (DDX18), mRNA
NM_030655	Homo sapiens DEAD/H (Asp-Glu-Ala-Asp/His) box polypeptide 11 (CHL1-like
	helicase homolog, S. cerevisiae) (DDX11), transcript variant 3, mRNA
NM_030653	Homo sapiens DEAD/H (Asp-Glu-Ala-Asp/His) box polypeptide 11 (CHL1-like
_	helicase homolog, S. cerevisiae) (DDX11), transcript variant 1, mRNA
NM_000770	Homo sapiens cytochrome P450, subfamily IIC (mephenytoin 4-hydroxylase),
	polypeptide 8 (CYP2C8), transcript variant Hp1-1, mRNA
NM_030878	Homo sapiens cytochrome P450, subfamily IIC (mephenytoin 4-hydroxylase),
	polypeptide 8 (CYP2C8), transcript variant Hp1-2, mRNA
NM_012239	Homo sapiens sirtuin silent mating type information regulation 2 homolog 3 (S.
-	cerevisiae) (SIRT3), mRNA
NM_030593	Homo sapiens sirtuin silent mating type information regulation 2 homolog 2 (S.
_	cerevisiae) (SIRT2), transcript variant 2, mRNA
NM_012237	Homo sapiens sirtuin silent mating type information regulation 2 homolog 2 (S.
_	cerevisiae) (SIRT2), transcript variant 1, mRNA
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NM_012238	Homo sapiens sirtuin silent mating type information regulation 2 homolog 1 (S. cerevisiae) (SIRT1), mRNA
NM_031309	Homo sapiens scratch homolog 1, zinc finger protein (Drosophila) (SCRT1),
_	mRNA
NM 031278	Homo sapiens tudor domain containing 1 (TDRD1), mRNA
NM 031277	Homo sapiens ring finger protein 17 (RNF17), transcript variant long, mRNA
NM 031276	Homo sapiens testis expressed sequence 11 (TEX11), mRNA
NM 031273	Homo sapiens testis expressed sequence 13B (TEX13B), mRNA
NM 031272	Homo sapiens testis expressed sequence 14 (TEX14), mRNA
NM 006636	Homo sapiens methylene tetrahydrofolate dehydrogenase (NAD+ dependent),
	methenyltetrahydrofolate cyclohydrolase (MTHFD2), nuclear gene encoding
	mitochondrial protein, mRNA
NM 022818	Homo sapiens microtubule-associated proteins 1A/1B light chain 3
-	(MAP1A/1BLC3), mRNA
NM 018607	Homo sapiens hypothetical protein PRO1853 (PRO1853), mRNA
NM 004856	Homo sapiens kinesin-like 5 (mitotic kinesin-like protein 1) (KNSL5), mRNA
NM 030979	Homo sapiens poly(A) binding protein, cytoplasmic 3 (PABPC3), mRNA
NM 030770	Homo sapiens transmembrane protease, serine 5 (spinesin) (TMPRSS5), mRNA
NM 002545	Homo sapiens opioid binding protein/cell adhesion molecule-like (OPCML),
1117_002545	mRNA
NM_014676	Homo sapiens pumilio homolog 1 (Drosophila) (PUM1), mRNA
NM 030673	Homo sapiens SEC13-like 1 (S. cerevisiae) (SEC13L1), mRNA
NM_003342	Homo sapiens ubiquitin-conjugating enzyme E2G 1 (UBC7 homolog, C. elegans) (UBE2G1), mRNA
NM 022051	Homo sapiens egl nine homolog 1 (C. elegans) (EGLN1), mRNA
NM 015577	Homo sapiens retinoic acid induced 14 (RAI14), mRNA
NM 012170	Homo sapiens F-box only protein 22 (FBXO22), mRNA
NM 022304	Homo sapiens histamine receptor H2 (HRH2), mRNA
NM_022333	Homo sapiens TIA1 cytotoxic granule-associated RNA binding protein-like 1 (TIAL1), transcript variant 2, mRNA
NM 003252	Homo sapiens TIA1 cytotoxic granule-associated RNA binding protein-like 1
14141_003232	(TIAL1), transcript variant 1, mRNA
NM 017910	Homo sapiens hypothetical protein FLJ20628 (FLJ20628), mRNA
NM 012384	Homo sapiens glucocorticoid modulatory element binding protein 2 (GMEB2),
	mRNA
NM_006118	Homo sapiens HS1 binding protein (HAX1), mRNA
NM_022740	Homo sapiens homeodomain interacting protein kinase 2 (HIPK2), mRNA
NM_002005	Homo sapiens feline sarcoma oncogene (FES), mRNA
NM_014757	Homo sapiens mastermind-like 1 (Drosophila) (MAML1), mRNA
NM_025136	Homo sapiens optic atrophy 3 (autosomal recessive, with chorea and spastic paraplegia) (OPA3), mRNA
NM_024505	Homo sapiens NADPH oxidase, EF hand calcium-binding domain 5 (NOX5), mRNA
NM_022362	Homo sapiens MMS19-like (MET18 homolog, S. cerevisiae) (MMS19L), mRNA
NM 000256	Homo sapiens myosin binding protein C, cardiac (MYBPC3), mRNA
	Homo sapiens oculocerebrorenal syndrome of Lowe (OCRL), transcript variant
NM_000276	a, mRNA
NM_001587	Homo sapiens oculocerebrorenal syndrome of Lowe (OCRL), transcript variant b, mRNA
NM 001407	Homo sapiens cadherin, EGF LAG seven-pass G-type receptor 3 (flamingo
	homolog, Drosophila) (CELSR3), mRNA

NM_001408	Homo sapiens cadherin, EGF LAG seven-pass G-type receptor 2 (flamingo
	homolog, Drosophila) (CELSR2), mRNA
NM_005735	Homo sapiens ARP1 actin-related protein 1 homolog B, centractin beta (yeast) (ACTR1B), mRNA
NM_012254	Homo sapiens very long-chain acyl-CoA synthetase homolog 2 (VLCS-H2), mRNA
NM_012331	Homo sapiens methionine sulfoxide reductase A (MSRA), mRNA
NM 016596	Homo sapiens histone deacetylase 7A (HDAC7A), transcript variant 2, mRNA
NM_015401	Homo sapiens histone deacetylase 7A (HDAC7A), transcript variant 1, mRNA
NM_004082	Homo sapiens dynactin 1 (p150, glued homolog, Drosophila) (DCTN1), transcript variant 1, mRNA
NM_023019	Homo sapiens dynactin 1 (p150, glued homolog, Drosophila) (DCTN1), transcript variant 2, mRNA
NM_002893	Homo sapiens retinoblastoma binding protein 7 (RBBP7), mRNA
NM_023001	Homo sapiens retinoblastoma binding protein 1 (RBBP1), transcript variant 3, mRNA
NM_023000	Homo sapiens retinoblastoma binding protein 1 (RBBP1), transcript variant 2, mRNA
NM_002892	Homo sapiens retinoblastoma binding protein 1 (RBBP1), transcript variant 1, mRNA
NM_024408	Homo sapiens Notch homolog 2 (Drosophila) (NOTCH2), mRNA
NM_012311	Homo sapiens KIN, antigenic determinant of recA protein homolog (mouse) (KIN), mRNA
NM_021938	Homo sapiens bruno-like 5, RNA binding protein (Drosophila) (BRUNOL5), mRNA
NM_020180	Homo sapiens bruno-like 4, RNA binding protein (Drosophila) (BRUNOL4), mRNA
NM_005868	Homo sapiens BET1 homolog (S. cerevisiae) (BET1), mRNA
NM_002467	Homo sapiens v-myc myelocytomatosis viral oncogene homolog (avian) (MYC), mRNA
NM_022817	Homo sapiens period homolog 2 (Drosophila) (PER2), transcript variant 1, mRNA
NM_003894	Homo sapiens period homolog 2 (Drosophila) (PER2), transcript variant 2, mRNA
NM_006660	Homo sapiens ClpX caseinolytic protease X homolog (E. coli) (CLPX), mRNA
NM_012394	Homo sapiens prefoldin 2 (PFDN2), mRNA
NM_004234	Homo sapiens zinc finger protein 93 homolog (mouse) (ZFP93), mRNA
NM_005870	Homo sapiens sin3-associated polypeptide, 18kD (SAP18), mRNA
NM_003350	Homo sapiens ubiquitin-conjugating enzyme E2 variant 2 (UBE2V2), mRNA
NM_022476	Homo sapiens fused toes homolog (mouse) (FTS), mRNA
NM_022444	Homo sapiens solute carrier family 13 (sodium/sulfate symporters), member 1 (SLC13A1), mRNA
NM_018127	Homo sapiens elaC homolog 2 (E. coli) (ELAC2), mRNA
NM_014317	Homo sapiens trans-prenyltransferase (TPT), mRNA
NM_022173	Homo sapiens TIA1 cytotoxic granule-associated RNA binding protein (TIA1), transcript variant 2, mRNA
NM_022037	Homo sapiens TIA1 cytotoxic granule-associated RNA binding protein (TIA1), transcript variant 1, mRNA
NM_004973	Homo sapiens jumonji homolog (mouse) (JMJ), mRNA
NM_021971	Homo sapiens GDP-mannose pyrophosphorylase B (GMPPB), transcript variant 2, mRNA
NM_013334	Homo sapiens GDP-mannose pyrophosphorylase B (GMPPB), transcript variant

	1, mRNA
NM_013335	Homo sapiens GDP-mannose pyrophosphorylase A (GMPPA), mRNA
NM_021267	Homo sapiens LAG1 longevity assurance homolog 1 (S. cerevisiae) (LASS1),
	mRNA
NM_005811	Homo sapiens growth differentiation factor 11 (GDF11), mRNA
NM_005971	Homo sapiens FXYD domain-containing ion transport regulator 3 (FXYD3), transcript variant 1, mRNA
NM 021910	Homo sapiens FXYD domain-containing ion transport regulator 3 (FXYD3),
	transcript variant 2, mRNA
NM 022096	Homo sapiens ankyrin repeat domain 5 (ANKRD5), mRNA
NM 022073	Homo sapiens egl nine homolog 3 (C. elegans) (EGLN3), mRNA
NM 022047	Homo sapiens differentially expressed in FDCP 6 homolog (mouse) (DEF6),
	mRNA
NM_021778	Homo sapiens a disintegrin and metalloproteinase domain 28 (ADAM28),
	transcript variant 2, mRNA
NM_021777	Homo sapiens a disintegrin and metalloproteinase domain 28 (ADAM28), transcript variant 3, mRNA
NM 000152	Homo sapiens glucosidase, alpha; acid (Pompe disease, glycogen storage disease
_	type II) (GAA), mRNA
NM_002910	Homo sapiens renin binding protein (RENBP), mRNA
NM_012072	Homo sapiens complement component 1, q subcomponent, receptor 1 (C1QR1), mRNA
NM_000534	Homo sapiens PMS1 postmeiotic segregation increased 1 (S. cerevisiae) (PMS1), mRNA
NM 005451	Homo sapiens enigma (LIM domain protein) (ENIGMA), mRNA
NM_021975	Homo sapiens v-rel reticuloendotheliosis viral oncogene homolog A, nuclear
1414_021575	factor of kappa light polypeptide gene enhancer in B-cells 3, p65 (avian)
	(RELA), mRNA
NM_021958	Homo sapiens H2.0-like homeo box 1 (Drosophila) (HLX1), mRNA
NM_004139	Homo sapiens lipopolysaccharide binding protein (LBP), mRNA
NM_005442	Homo sapiens eomesodermin homolog (Xenopus laevis) (EOMES), mRNA
NM_004187	Homo sapiens Smcx homolog, X chromosome (mouse) (SMCX), mRNA
NM_003170	Homo sapiens suppressor of Ty 6 homolog (S. cerevisiae) (SUPT6H), mRNA
NM_003062	Homo sapiens slit homolog 3 (Drosophila) (SLIT3), mRNA
NM_003068	Homo sapiens slug homolog, zinc finger protein (chicken) (SLUG), mRNA
NM_021824	Homo sapiens NIF3 NGG1 interacting factor 3-like 1 (S. pombe) (NIF3L1), mRNA
NM 021783	Homo sapiens ectodysplasin A2 isoform receptor (XEDAR), mRNA
NM_004196	Homo sapiens cyclin-dependent kinase-like 1 (CDC2-related kinase) (CDKL1), mRNA
NM_000535	Homo sapiens PMS2 postmeiotic segregation increased 2 (S. cerevisiae) (PMS2),
	mRNA
NM_002356	Homo sapiens myristoylated alanine-rich protein kinase C substrate (MARCKS), mRNA
NM 021728	Homo sapiens orthodenticle homolog 2 (Drosophila) (OTX2), mRNA
NM_014588	Homo sapiens visual system homeobox 1 homolog, CHX10-like (zebrafish) (VSX1), mRNA
NM 003503	Homo sapiens CDC7 cell division cycle 7-like 1 (S. cerevisiae) (CDC7L1),
TATAT_002202	mRNA
NM_004059	Homo sapiens cysteine conjugate-beta lyase; cytoplasmic (glutamine.
	transaminase K, kyneurenine aminotransferase) (CCBL1), mRNA
NM 020651	Homo sapiens pellino homolog 1 (Drosophila) (PELI1), mRNA

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NM_018411	Homo sapiens hairless homolog (mouse) (HR), mRNA
NM_014569	Homo sapiens zinc finger protein 95 homolog (mouse) (ZFP95), mRNA
NM_012458	Homo sapiens translocase of inner mitochondrial membrane 13 homolog B (yeast) (TIMM13B), mRNA
NM_000672	Homo sapiens alcohol dehydrogenase 6 (class V) (ADH6), mRNA
NM_003603	Homo sapiens Arg/Abl-interacting protein ArgBP2 (ARGBP2), transcript variant 1, mRNA
NM_021069	Homo sapiens Arg/Abl-interacting protein ArgBP2 (ARGBP2), transcript variant 2, mRNA
NM_004950	Homo sapiens dermatan sulfate proteoglycan 3 (DSPG3), mRNA
NM_004701	Homo sapiens cyclin B2 (CCNB2), mRNA
NM_021100	Homo sapiens NFS1 nitrogen fixation 1 (S. cerevisiae) (NFS1), mRNA
NM_021255	Homo sapiens pellino homolog 2 (Drosophila) (PELI2), mRNA
NM_021115	Homo sapiens seizure related 6 homolog (mouse)-like (SEZ6L), mRNA
NM_004756	Homo sapiens numb homolog (Drosophila)-like (NUMBL), mRNA
NM_004690	Homo sapiens LATS, large tumor suppressor, homolog 1 (Drosophila) (LATS1), mRNA
NM_000461	Homo sapiens thyroid hormone receptor, beta (erythroblastic leukemia viral (verb-a) oncogene homolog 2, avian) (THRB), mRNA
NM_021078	Homo sapiens GCN5 general control of amino-acid synthesis 5-like 2 (yeast) (GCN5L2), mRNA
NM_002877	Homo sapiens RAD51-like 1 (S. cerevisiae) (RAD51L1), mRNA
NM_001552	Homo sapiens insulin-like growth factor binding protein 4 (IGFBP4), mRNA
NM_002487	Homo sapiens necdin homolog (mouse) (NDN), mRNA
NM_012425	Homo sapiens Ras suppressor protein 1 (RSU1), mRNA
NM_005618	Homo sapiens delta-like 1 (Drosophila) (DLL1), mRNA
NM_021038	Homo sapiens muscleblind-like (Drosophila) (MBNL), mRNA
NM_014268	Homo sapiens microtubule-associated protein, RP/EB family, member 2 (MAPRE2), mRNA
NM_020662	Homo sapiens MRS2-like, magnesium homeostasis factor (S. cerevisiae) (MRS2L), mRNA
NM_020649	Homo sapiens chromobox homolog 8 (Pc class homolog, Drosophila) (CBX8), mRNA
NM_018436	Homo sapiens allantoicase (ALLC), mRNA
NM_020528	Homo sapiens poly(rC) binding protein 3 (PCBP3), mRNA
NM_014276	Homo sapiens recombining binding protein suppressor of hairless (Drosophila)-like (RBPSUHL), mRNA
NM_019557	Homo sapiens hypothetical protein RP1-317E23 (LOC56181), mRNA
NM_020347	Homo sapiens leucine zipper transcription factor-like 1 (LZTFL1), mRNA
NM_005744	Homo sapiens ariadne homolog, ubiquitin-conjugating enzyme E2 binding protein, 1 (Drosophila) (ARIH1), mRNA
NM_007044	Homo sapiens katanin p60 (ATPase-containing) subunit A 1 (KATNA1), mRNA
NM_002688	Homo sapiens peanut-like 1 (Drosophila) (PNUTL1), mRNA
NM_013384	Homo sapiens LAG1 longevity assurance homolog 2 (S. cerevisiae) (LASS2), mRNA
NM_020230	Homo sapiens peter pan homolog (Drosophila) (PPAN), mRNA
NM_020182	Homo sapiens transmembrane, prostate androgen induced RNA (TMEPAI), mRNA
NM_020248	Homo sapiens catenin, beta interacting protein 1 (CTNNBIP1), mRNA
NM_000399	Homo sapiens early growth response 2 (Krox-20 homolog, Drosophila) (EGR2), mRNA
NM_002965	Homo sapiens S100 calcium binding protein A9 (calgranulin B) (S100A9),

mRNA
Homo sapiens S100 calcium binding protein A8 (calgranulin A) (S100A8),
mRNA
Homo sapiens S100 calcium binding protein A7 (psoriasin 1) (S100A7), mRNA
Homo sapiens S100 calcium binding protein A6 (calcyclin) (S100A6), mRNA
Homo sapiens S100 calcium binding protein A4 (calcium protein, calvasculin,
metastasin, murine placental homolog) (S100A4), transcript variant 2, mRNA
Homo sapiens S100 calcium binding protein A4 (calcium protein, calvasculin,
metastasin, murine placental homolog) (S100A4), transcript variant 1, mRNA
Homo sapiens S100 calcium binding protein A2 (S100A2), mRNA
Homo sapiens ornithine decarboxylase antizyme 2 (OAZ2), mRNA
Homo sapiens HMT1 hnRNP methyltransferase-like 3 (S. cerevisiae)
(HRMT1L3), mRNA
Homo sapiens par-3 partitioning defective 3 homolog (C. elegans) (PARD3), mRNA
Homo sapiens staufen, RNA binding protein (Drosophila) (STAU), transcript variant T1, mRNA
Homo sapiens staufen, RNA binding protein (Drosophila) (STAU), transcript
variant T3, mRNA
Homo sapiens staufen, RNA binding protein (Drosophila) (STAU), transcript
variant T2, mRNA
Homo sapiens G antigen, family B, 1 (prostate associated) (GAGEB1), mRNA
Homo sapiens golgi associated, gamma adaptin ear containing, ARF binding
protein 2 (GGA2), mRNA
Homo sapiens golgi associated, gamma adaptin ear containing, ARF binding
protein 1 (GGA1), mRNA
Homo sapiens vesicle-associated membrane protein 3 (cellubrevin) (VAMP3), mRNA
Homo sapiens anillin, actin binding protein (scraps homolog, Drosophila)
(ANLN), mRNA
Homo sapiens mitofusin 1 (MFN1), transcript variant 2, mRNA
Homo sapiens spermatid perinuclear RNA binding protein (STRBP), mRNA
Homo sapiens F-box and leucine-rich repeat protein 8 (FBXL8), mRNA
Homo sapiens solute carrier family 4 (anion exchanger), member 1, adaptor
protein (SLC4A1AP), mRNA
Homo sapiens LUC7-like (S. cerevisiae) (LUC7L), mRNA
Homo sapiens chromosome 17 open reading frame 31 (C17orf31), mRNA
Homo sapiens elaC homolog 1 (E. coli) (ELAC1), mRNA
Homo sapiens activated p21cdc42Hs kinase (ACK1), mRNA
Homo sapiens period homolog 3 (Drosophila) (PER3), mRNA
Homo sapiens Wiskott-Aldrich syndrome protein interacting protein (WASPIP), mRNA
Homo sapiens tubulin-specific chaperone d (TBCD), mRNA
Homo sapiens secreted frizzled-related protein 4 (SFRP4), mRNA
Homo sapiens retinol binding protein 4, plasma (RBP4), mRNA
Homo sapiens retinol binding protein 1, cellular (RBP1), mRNA
Homo sapiens hairy homolog (Drosophila) (HRY), mRNA
Homo sapiens v-crk sarcoma virus CT10 oncogene homolog (avian) (CRK),
ranscript variant I, mRNA
Homo sapiens v-crk sarcoma virus CT10 oncogene homolog (avian) (CRK), ranscript variant II, mRNA
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	(PARD6A), mRNA
NM 017420	Homo sapiens sine oculis homeobox homolog 4 (Drosophila) (SIX4), mRNA
NM 016932	Homo sapiens sine oculis homeobox homolog 4 (Drosophila) (SIX4), mRNA
NM_017415	Homo sapiens kelch-like 3 (Drosophila) (KLHL3), mRNA
NM_017412	Homo sapiens frizzled homolog 3 (Drosophila) (FZD3), mRNA
NM_003400	Homo sapiens exportin 1 (CRM1 homolog, yeast) (XPO1), mRNA
NM_002889	Homo sapiens retinoic acid receptor responder (tazarotene induced) 2 (RARRES2), mRNA
NM_006064	Homo sapiens GTP-binding protein ragB (RAGB), transcript variant RAGBs, mRNA
NM_016656	Homo sapiens GTP-binding protein ragB (RAGB), transcript variant RAGBl, mRNA
NM 003857	Homo sapiens galanin receptor 2 (GALR2), mRNA
NM_016655	Homo sapiens GA binding protein transcription factor, beta subunit 2 (47kD) (GABPB2), transcript variant gamma, mRNA
NM_002041	Homo sapiens GA binding protein transcription factor, beta subunit 2 (47kD) (GABPB2), transcript variant gamma, mRNA
NM_016654	Homo sapiens GA binding protein transcription factor, beta subunit 1 (53kD) (GABPB1), transcript variant beta, mRNA
NM_005254	Homo sapiens GA binding protein transcription factor, beta subunit 1 (53kD) (GABPB1), transcript variant beta, mRNA
NM_015843	Homo sapiens LIM domain only 7 (LMO7), transcript variant 3, mRNA
NM 015842	Homo sapiens LIM domain only 7 (LMO7), transcript variant 2, mRNA
NM 002228	Homo sapiens v-jun sarcoma virus 17 oncogene homolog (avian) (JUN), mRNA
NM 016178	Homo sapiens ornithine decarboxylase antizyme 3 (OAZ3), mRNA
NM_016538	Homo sapiens sirtuin silent mating type information regulation 2 homolog 7 (S.
	cerevisiae) (SIRT7), mRNA
NM_016539	Homo sapiens sirtuin silent mating type information regulation 2 homolog 6 (S. cerevisiae) (SIRT6), mRNA
NM 016316	Homo sapiens REV1-like (yeast) (REV1L), mRNA
NM_016138	Homo sapiens COQ7 coenzyme Q, 7 homolog ubiquinone (yeast) (COQ7), mRNA
NM_016583	Homo sapiens palate, lung and nasal epithelium carcinoma associated (PLUNC), mRNA
NM 015886	Homo sapiens protease inhibitor 15 (PI15), mRNA
NM_016067	Homo sapiens mitochondrial ribosomal protein S18C (MRPS18C), nuclear gene encoding mitochondrial protein, mRNA
NM 015946	Homo sapiens pelota homolog (Drosophila) (PELO), mRNA
NM 016397	Homo sapiens TH1-like (Drosophila) (TH1L), mRNA
NM_016587	Homo sapiens chromobox homolog 3 (HP1 gamma homolog, Drosophila) (CBX3), mRNA
NM 016347	Homo sapiens putative N-acetyltransferase Camello 2 (CML2), mRNA
NM 015727	Homo sapiens tachykinin receptor 1 (TACR1), transcript variant short, mRNA
NM 001058	Homo sapiens tachykinin receptor 1 (TACR1), transcript variant long, mRNA
NM 004052	Homo sapiens BCL2/adenovirus E1B 19kD interacting protein 3 (BNIP3),
	nuclear gene encoding mitochondrial protein, mRNA
NM_014820	Homo sapiens translocase of outer mitochondrial membrane 70 homolog A (yeast) (TOMM70A), mRNA
NM 014918	Homo sapiens carbohydrate (chondroitin) synthase 1 (CHSY1), mRNA
NM_014707	Homo sapiens histone deacetylase 9 (HDAC9-PENDING), transcript variant 3, mRNA
NM 014683	Homo sapiens unc-51-like kinase 2 (C. elegans) (ULK2), mRNA
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NM_014874	Homo sapiens mitofusin 2 (MFN2), mRNA
NM_014071	Homo sapiens nuclear receptor coactivator 6 (NCOA6), mRNA
NM_015700	Homo sapiens HIRA interacting protein 5 (HIRIP5), mRNA
NM_015685	Homo sapiens syndecan binding protein (syntenin) 2 (SDCBP2), mRNA
NM_014263	Homo sapiens YME1-like 1 (S. cerevisiae) (YME1L1), mRNA
NM_014297	Homo sapiens protein expressed in thyroid (YF13H12), mRNA
NM_014393	Homo sapiens staufen, RNA binding protein, homolog 2 (Drosophila) (STAU2), mRNA
NM_014403	Homo sapiens sialyltransferase 7D ((alpha-N-acetylneuraminyl-2,3-beta-galactosyl-1,3)-N-acetyl galactosaminide alpha-2,6-sialyltransferase) (SIAT7D), mRNA
NM_014465	Homo sapiens sulfotransferase family, cytosolic, 1B, member 1 (SULT1B1), mRNA
NM_014485	Homo sapiens prostaglandin D2 synthase, hematopoietic (PGDS), mRNA
NM_014303	Homo sapiens pescadillo homolog 1, containing BRCT domain (zebrafish) (PES1), mRNA
NM_014253	Homo sapiens odz, odd Oz/ten-m homolog 1(Drosophila) (ODZ1), mRNA
NM_014429	Homo sapiens microrchidia homolog (mouse) (MORC), mRNA
NM_006439	Homo sapiens mab-21-like 2 (C. elegans) (MAB21L2), mRNA
NM_015322	Homo sapiens fem-1 homolog b (C. elegans) (FEM1B), mRNA
NM_014591	Homo sapiens Kv channel interacting protein 2 (KCNIP2), mRNA
NM_004449	Homo sapiens v-ets erythroblastosis virus E26 oncogene like (avian) (ERG), mRNA
NM_014420	Homo sapiens dickkopf homolog 4 (Xenopus laevis) (DKK4), mRNA
NM_014421	Homo sapiens dickkopf homolog 2 (Xenopus laevis) (DKK2), mRNA
NM_014325	Homo sapiens coronin, actin binding protein, 1C (CORO1C), mRNA
NM_014246	Homo sapiens cadherin, EGF LAG seven-pass G-type receptor 1 (flamingo homolog, Drosophila) (CELSR1), mRNA
NM_014391	Homo sapiens cardiac ankyrin repeat protein (CARP), mRNA
NM_014336	Homo sapiens aryl hydrocarbon receptor interacting protein-like 1 (AIPL1), mRNA
NM_014265	Homo sapiens a disintegrin and metalloproteinase domain 28 (ADAM28), transcript variant 1, mRNA
NM_014237	Homo sapiens a disintegrin and metalloproteinase domain 18 (ADAM18), mRNA
NM_005032	Homo sapiens plastin 3 (T isoform) (PLS3), mRNA
NM_013980	Homo sapiens BCL2/adenovirus E1B 19kD interacting protein 1 (BNIP1), transcript variant BNIP1-c, mRNA
NM_013979	Homo sapiens BCL2/adenovirus E1B 19kD interacting protein 1 (BNIP1), transcript variant BNIP1-b, mRNA
NM_013978	Homo sapiens BCL2/adenovirus E1B 19kD interacting protein 1 (BNIP1), transcript variant BNIP1-a, mRNA
NM_004178	Homo sapiens TAR (HIV) RNA binding protein 2 (TARBP2), mRNA
NM_005915	Homo sapiens MCM6 minichromosome maintenance deficient 6 (MIS5 homolog, S. pombe) (S. cerevisiae) (MCM6), mRNA
NM_002576	Homo sapiens p21/Cdc42/Rac1-activated kinase 1 (STE20 homolog, yeast) (PAK1), mRNA
NM_012091	Homo sapiens adenosine deaminase, tRNA-specific 1 (ADAT1), mRNA
NM_005358	Homo sapiens LIM domain only 7 (LMO7), mRNA
NM_013451	Homo sapiens fer-1-like 3, myoferlin (C. elegans) (FER1L3), mRNA
NM_006113	Homo sapiens vav 3 oncogene (VAV3), mRNA
NM_003869	Homo sapiens carboxylesterase 2 (intestine, liver) (CES2), mRNA
	1 Colorado L (modello, nvol) (Colon), matrix

NM_005721	Homo sapiens ARP3 actin-related protein 3 homolog (yeast) (ACTR3), mRNA
NM_003325	Homo sapiens HIR histone cell cycle regulation defective homolog A (S.
	cerevisiae) (HIRA), mRNA
NM_012242	Homo sapiens dickkopf homolog 1 (Xenopus laevis) (DKK1), mRNA
NM_012429	Homo sapiens SEC14-like 2 (S. cerevisiae) (SEC14L2), mRNA
NM_012190	Homo sapiens formyltetrahydrofolate dehydrogenase (FTHFD), mRNA
NM_005069	Homo sapiens single-minded homolog 2 (Drosophila) (SIM2), transcript variant SIM2, mRNA
NM_009586	Homo sapiens single-minded homolog 2 (Drosophila) (SIM2), transcript variant SIM2s, mRNA
NM_002610	Homo sapiens pyruvate dehydrogenase kinase, isoenzyme 1 (PDK1), nuclear gene encoding mitochondrial protein, mRNA
NM 013374	Homo sapiens programmed cell death 6 interacting protein (PDCD6IP), mRNA
NM 013367	Homo sapiens anaphase-promoting complex subunit 4 (APC4), mRNA
NM 002968	Homo sapiens sal-like 1 (Drosophila) (SALL1), mRNA
NM 002449	Homo sapiens msh homeo box homolog 2 (Drosophila) (MSX2), mRNA
NM_006739	Homo sapiens MCM5 minichromosome maintenance deficient 5, cell division
_	cycle 46 (S. cerevisiae) (MCM5), mRNA
NM_012460	Homo sapiens translocase of inner mitochondrial membrane 9 homolog (yeast) (TIMM9), mRNA
NM_012457	Homo sapiens translocase of inner mitochondrial membrane 13 homolog A (yeast) (TIMM13A), mRNA
NM_012456	Homo sapiens translocase of inner mitochondrial membrane 10 homolog (yeast) (TIMM10), mRNA
NM_012450	Homo sapiens solute carrier family 13 (sodium/sulfate symporters), member 4 (SLC13A4), mRNA
NM_012444	Homo sapiens SPO11 meiotic protein covalently bound to DSB-like (S. cerevisiae) (SPO11), mRNA
NM_012240	Homo sapiens sirtuin silent mating type information regulation 2 homolog 4 (S. cerevisiae) (SIRT4), mRNA
NM_012387	Homo sapiens peptidyl arginine deiminase, type V (PAD), mRNA
NM_012381	Homo sapiens origin recognition complex, subunit 3-like (yeast) (ORC3L), mRNA
NM_012225	Homo sapiens nucleotide binding protein 2 (MinD homolog, E. coli) (NUBP2), mRNA
NM 012222	Homo sapiens mutY homolog (E. coli) (MUTYH), mRNA
NM_012279	Homo sapiens double-stranded RNA-binding zinc finger protein JAZ (JAZ), mRNA
NM_012206	Homo sapiens hepatitis A virus cellular receptor 1 (HAVCR-1), mRNA
NM 012205	Homo sapiens 3-hydroxyanthranilate 3,4-dioxygenase (HAAO), mRNA
NM 012198	Homo sapiens grancalcin, EF-hand calcium binding protein (GCA), mRNA
NM 012193	Homo sapiens frizzled homolog 4 (Drosophila) (FZD4), mRNA
NM 012192	Homo sapiens fracture callus 1 homolog (rat) (FXC1), mRNA
NM 012076	Homo sapiens crumbs homolog 1 (Drosophila) (CRB1), mRNA
NM_012124	Homo sapiens cysteine and histidine-rich domain (CHORD)-containing, zinc binding protein 1 (CHORDC1), mRNA
NM_012118	Homo sapiens CCR4 carbon catabolite repression 4-like (S. cerevisiae) (CCRN4L), mRNA
NM_012117	Homo sapiens chromobox homolog 5 (HP1 alpha homolog, Drosophila) (CBX5), mRNA
NM 012108	Homo sapiens BCR downstream signaling 1 (BRDG1), mRNA
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NM_012094	Homo sapiens peroxiredoxin 5 (PRDX5), mRNA
NM_004506	Homo sapiens heat shock transcription factor 2 (HSF2), mRNA
NM_004423	Homo sapiens dishevelled, dsh homolog 3 (Drosophila) (DVL3), mRNA
NM_007374	Homo sapiens sine oculis homeobox homolog 6 (Drosophila (SIX6), mRNA
NM_007373	Homo sapiens soc-2 suppressor of clear homolog (C. elegans) (SHOC2), mRNA
NM_002388	Homo sapiens MCM3 minichromosome maintenance deficient 3 (S. cerevisiae) (MCM3), mRNA
NM_004873	Homo sapiens BCL2-associated athanogene 5 (BAG5), mRNA
NM_007316	Homo sapiens agouti related protein homolog (mouse) (AGRP), transcript variant 2, mRNA
NM_003819	Homo sapiens poly(A) binding protein, cytoplasmic 4 (inducible form) (PABPC4), mRNA
NM 005737	Homo sapiens ADP-ribosylation factor-like 7 (ARL7), mRNA
NM 002358	Homo sapiens MAD2 mitotic arrest deficient-like 1 (yeast) (MAD2L1), mRNA
NM 007264	Homo sapiens adrenomedullin receptor (ADMR), mRNA
NM 006870	Homo sapiens destrin (actin depolymerizing factor) (DSTN), mRNA
NM_005476	Homo sapiens UDP-N-acetylglucosamine-2-epimerase/N-acetylmannosamine kinase (GNE), mRNA
NM_007309	Homo sapiens diaphanous homolog 2 (Drosophila) (DIAPH2), transcript variant 12C, mRNA
NM 001878	Homo sapiens cellular retinoic acid binding protein 2 (CRABP2), mRNA
NM_000489	Homo sapiens alpha thalassemia/mental retardation syndrome X-linked (RAD54 homolog, S. cerevisiae) (ATRX), mRNA
NM 002528	Homo sapiens nth endonuclease III-like 1 (E. coli) (NTHL1), mRNA
NM_004085	Homo sapiens translocase of inner mitochondrial membrane 8 homolog A (yeast)
11112_00 1000	(TIMM8A), nuclear gene encoding mitochondrial protein, mRNA
NM 002310	Homo sapiens leukemia inhibitory factor receptor (LIFR), mRNA
NM 004733	Homo sapiens acetyl-Coenzyme A transporter (ACATN), mRNA
NM 002657	Homo sapiens pleiomorphic adenoma gene-like 2 (PLAGL2), mRNA
NM 006724	Homo sapiens mitogen-activated protein kinase kinase kinase 4 (MAP3K4),
	transcript variant 2, mRNA
NM 006882	Homo sapiens Mdm2, transformed 3T3 cell double minute 2, p53 binding protein
	(mouse) (MDM2), transcript variant MDM2e, mRNA
NM_006881	Homo sapiens Mdm2, transformed 3T3 cell double minute 2, p53 binding protein
	(mouse) (MDM2), transcript variant MDM2d, mRNA
NM_006880	Homo sapiens Mdm2, transformed 3T3 cell double minute 2, p53 binding protein
_	(mouse) (MDM2), transcript variant MDM2c, mRNA
NM_006879	Homo sapiens Mdm2, transformed 3T3 cell double minute 2, p53 binding protein
	(mouse) (MDM2), transcript variant MDM2b, mRNA
NM_006878	Homo sapiens Mdm2, transformed 3T3 cell double minute 2, p53 binding protein
	(mouse) (MDM2), transcript variant MDM2a, mRNA
NM_003801	Homo sapiens GPAA1P anchor attachment protein 1 homolog (yeast) (GPAA1), mRNA
NM_003193	Homo sapiens tubulin-specific chaperone e (TBCE), mRNA
NM 002370	Homo sapiens mago-nashi homolog, proliferation-associated (Drosophila)
	(MAGOH), mRNA
NM 006341	Homo sapiens MAD2 mitotic arrest deficient-like 2 (yeast) (MAD2L2), mRNA
NM 006149	Homo sapiens lectin, galactoside-binding, soluble, 4 (galectin 4) (LGALS4),
	mRNA
NM 003585	Homo sapiens double C2-like domains, beta (DOC2B), mRNA
NM_007129	Homo sapiens Zic family member 2 (odd-paired homolog, Drosophila) (ZIC2),
	mRNA
L	IIIANA

NM 007279	Homo sapiens U2 small nuclear ribonucleoprotein auxiliary factor (65kD)
_	(U2AF65), mRNA
NM_007194	Homo sapiens CHK2 checkpoint homolog (S. pombe) (CHEK2), mRNA
NM_007271	Homo sapiens serine/threonine kinase 38 (STK38), mRNA
NM_007232	Homo sapiens histamine receptor H3 (HRH3), mRNA
NM_007278	Homo sapiens GABA(A) receptor-associated protein (GABARAP), mRNA
NM_007197	Homo sapiens frizzled homolog 10 (Drosophila) (FZD10), mRNA
NM_007246	Homo sapiens kelch-like 2, Mayven (Drosophila) (KLHL2), mRNA
NM_001466	Homo sapiens frizzled homolog 2 (Drosophila) (FZD2), mRNA
NM_006482	Homo sapiens dual-specificity tyrosine-(Y)-phosphorylation regulated kinase 2 (DYRK2), transcript variant 2, mRNA
NM_003583	Homo sapiens dual-specificity tyrosine-(Y)-phosphorylation regulated kinase 2 (DYRK2), transcript variant 1, mRNA
NM_006484	Homo sapiens dual-specificity tyrosine-(Y)-phosphorylation regulated kinase 1B (DYRK1B), transcript variant c, mRNA
NM_006483	Homo sapiens dual-specificity tyrosine-(Y)-phosphorylation regulated kinase 1B (DYRK1B), transcript variant b, mRNA
NM_001882	Homo sapiens corticotropin releasing hormone binding protein (CRHBP), mRNA
NM_005889	Homo sapiens apolipoprotein B mRNA editing enzyme, catalytic polypeptide 1 (APOBEC1), transcript variant 2, mRNA
NM_001644	Homo sapiens apolipoprotein B mRNA editing enzyme, catalytic polypeptide 1 (APOBEC1), transcript variant 1, mRNA
NM_006936	Homo sapiens SMT3 suppressor of mif two 3 homolog 1 (yeast) (SMT3H1), mRNA
NM 006912	Homo sapiens Ric-like, expressed in many tissues (Drosophila) (RIT), mRNA
NM 006910	Homo sapiens retinoblastoma binding protein 6 (RBBP6), mRNA
NM_007068	Homo sapiens DMC1 dosage suppressor of mck1 homolog, meiosis-specific homologous recombination (yeast) (DMC1), mRNA
NM 007021	Homo sapiens decidual protein induced by progesterone (DEPP), mRNA
NM_007007	Homo sapiens cleavage and polyadenylation specific factor 6, 68kD subunit (CPSF6), mRNA
NM_006822	Homo sapiens GTP-binding protein homologous to Saccharomyces cerevisiae SEC4 (SEC4L), mRNA
NM_006843	Homo sapiens serine dehydratase (SDS), mRNA
NM_006746	Homo sapiens sex comb on midleg-like 1 (Drosophila) (SCML1), mRNA
NM_006824	Homo sapiens EBNA1 binding protein 2 (EBNA1BP2), mRNA
NM_005922	Homo sapiens mitogen-activated protein kinase kinase kinase 4 (MAP3K4), transcript variant 1, mRNA
NM_006807	Homo sapiens chromobox homolog 1 (HP1 beta homolog Drosophila) (CBX1), mRNA
NM_006734	Homo sapiens human immunodeficiency virus type I enhancer binding protein 2 (HIVEP2), mRNA
NM_006732	Homo sapiens FBJ murine osteosarcoma viral oncogene homolog B (FOSB), mRNA
NM_006729	Homo sapiens diaphanous homolog 2 (Drosophila) (DIAPH2), transcript variant 156, mRNA
NM_006829	Homo sapiens adipose specific 2 (APM2), mRNA
NM_006872	Homo sapiens TFIIA-alpha/beta-like factor (ALF), mRNA
NM_006796	Homo sapiens AFG3 ATPase family gene 3-like 2 (yeast) (AFG3L2), nuclear gene encoding mitochondrial protein, mRNA
NM_006544	Homo sapiens SEC10-like 1 (S. cerevisiae) (SEC10L1), mRNA

NM_006666	Homo sapiens RuvB-like 2 (E. coli) (RUVBL2), mRNA
NM_006509	Homo sapiens v-rel reticuloendotheliosis viral oncogene homolog B, nuclear
	factor of kappa light polypeptide gene enhancer in B-cells 3 (avian) (RELB),
275.00660	mRNA
NM_006606	Homo sapiens retinoblastoma binding protein 9 (RBBP9), mRNA
NM_006620	Homo sapiens HBS1-like (S. cerevisiae) (HBS1L), mRNA
NM_006561	Homo sapiens CUG triplet repeat, RNA binding protein 2 (CUGBP2), mRNA
NM_006579	Homo sapiens emopamil binding protein (sterol isomerase) (EBP), mRNA
NM_006560	Homo sapiens CUG triplet repeat, RNA binding protein 1 (CUGBP1), mRNA
NM_001211	Homo sapiens BUB1 budding uninhibited by benzimidazoles 1 homolog beta (yeast) (BUB1B), mRNA
NM_006374	Homo sapiens serine/threonine kinase 25 (STE20 homolog, yeast) (STK25), mRNA
NM_006377	Homo sapiens unc-13-like (C. elegans) (UNC13), mRNA
NM_006357	Homo sapiens ubiquitin-conjugating enzyme E2E 3 (UBC4/5 homolog, yeast) (UBE2E3), mRNA
NM_006323	Homo sapiens SEC24 related gene family, member B (S. cerevisiae) (SEC24B), mRNA
NM_006364	Homo sapiens Sec23 homolog A (S. cerevisiae) (SEC23A), mRNA
NM_006272	Homo sapiens S100 calcium binding protein, beta (neural) (S100B), mRNA
NM_006271	Homo sapiens S100 calcium binding protein A1 (S100A1), mRNA
NM_006391	Homo sapiens RAN binding protein 7 (RANBP7), mRNA
NM_006265	Homo sapiens RAD21 homolog (S. pombe) (RAD21), mRNA
NM_006203	Homo sapiens phosphodiesterase 4D, cAMP-specific (phosphodiesterase E3
	dunce homolog, Drosophila) (PDE4D), mRNA
NM_006202	Homo sapiens phosphodiesterase 4A, cAMP-specific (phosphodiesterase E2 dunce homolog, Drosophila) (PDE4A), mRNA
NM_006190	Homo sapiens origin recognition complex, subunit 2-like (yeast) (ORC2L), mRNA
NM_006181	Homo sapiens netrin 2-like (chicken) (NTN2L), mRNA
NM_006168	Homo sapiens NK6 transcription factor homolog A (Drosophila) (NKX6A), mRNA
NM_006167	Homo sapiens NK3 transcription factor homolog A (Drosophila) (NKX3A), mRNA
NM_006159	Homo sapiens NEL-like 2 (chicken) (NELL2), mRNA
NM_006157	Homo sapiens NEL-like 1 (chicken) (NELL1), mRNA
NM_005360	Homo sapiens v-maf musculoaponeurotic fibrosarcoma oncogene homolog (avian) (MAF), mRNA
NM_006306	Homo sapiens SMC1 structural maintenance of chromosomes 1-like 1 (yeast) (SMC1L1), mRNA
NM_006461	Homo sapiens mitotic spindle coiled-coil related protein (DEEPEST), mRNA
NM_006314	Homo sapiens connector enhancer of KSR-like (Drosophila kinase suppressor of ras) (CNK1), mRNA
NM_006366	Homo sapiens adenylyl cyclase-associated protein 2 (CAP2), mRNA
NM_006444	Homo sapiens SMC2 structural maintenance of chromosomes 2-like 1 (yeast) (SMC2L1), mRNA
NM_006321	Homo sapiens ariadne homolog 2 (Drosophila) (ARIH2), mRNA
NM_006406	Homo sapiens peroxiredoxin 4 (PRDX4), mRNA
NM_006334	Homo sapiens olfactomedin 1 (OLFM1), transcript variant 2, mRNA
NM_004032	Homo sapiens D-aspartate oxidase (DDO), transcript variant 2, mRNA
NM_005985	Homo sapiens snail 1 homolog, zinc finger protein (Drosophila) (SNAI1), mRNA

	: GVD11 - 1 - (G 1 -) (GVD1) DNA
NM_006109	Homo sapiens SKB1 homolog (S. pombe) (SKB1), mRNA
NM_005982	Homo sapiens sine oculis homeobox homolog 1 (Drosophila) (SIX1), mRNA
NM_006089	Homo sapiens sex comb on midleg-like 2 (Drosophila) (SCML2), mRNA
NM_005980	Homo sapiens S100 calcium binding protein P (S100P), mRNA
NM_005979	Homo sapiens S100 calcium binding protein A13 (S100A13), mRNA
NM_005938	Homo sapiens myeloid/lymphoid or mixed-lineage leukemia (trithorax homolog, Drosophila); translocated to, 7 (MLLT7), mRNA
NM_005937	Homo sapiens myeloid/lymphoid or mixed-lineage leukemia (trithorax homolog, Drosophila); translocated to, 6 (MLLT6), mRNA
NM_005936	Homo sapiens myeloid/lymphoid or mixed-lineage leukemia (trithorax homolog, Drosophila); translocated to, 4 (MLLT4), mRNA
NM_005935	Homo sapiens myeloid/lymphoid or mixed-lineage leukemia (trithorax homolog, Drosophila); translocated to, 2 (MLLT2), mRNA
NM_005934	Homo sapiens myeloid/lymphoid or mixed-lineage leukemia (trithorax homolog, Drosophila); translocated to, 1 (MLLT1), mRNA
NM_005933	Homo sapiens myeloid/lymphoid or mixed-lineage leukemia (trithorax homolog, Drosophila) (MLL), mRNA
NM_005905	Homo sapiens MAD, mothers against decapentaplegic homolog 9 (Drosophila) (MADH9), mRNA
NM_005904	Homo sapiens MAD, mothers against decapentaplegic homolog 7 (Drosophila) (MADH7), mRNA
NM_005903	Homo sapiens MAD, mothers against decapentaplegic homolog 5 (Drosophila) (MADH5), mRNA
NM_005902	Homo sapiens MAD, mothers against decapentaplegic homolog 3 (Drosophila) (MADH3), mRNA
NM_005901	Homo sapiens MAD, mothers against decapentaplegic homolog 2 (Drosophila) (MADH2), mRNA
NM_005900	Homo sapiens MAD, mothers against decapentaplegic homolog 1 (Drosophila) (MADH1), mRNA
NM 006033	Homo sapiens lipase, endothelial (LIPG), mRNA
NM_006048	Homo sapiens ubiquitination factor E4B (UFD2 homolog, yeast) (UBE4B), mRNA
NM_006111	Homo sapiens acetyl-Coenzyme A acyltransferase 2 (mitochondrial 3-oxoacyl-Coenzyme A thiolase) (ACAA2), nuclear gene encoding mitochondrial protein, mRNA
NM_006012	Homo sapiens ClpP caseinolytic protease, ATP-dependent, proteolytic subunit homolog (E. coli) (CLPP), nuclear gene encoding mitochondrial protein, mRNA
NM_006110	Homo sapiens CD2 antigen (cytoplasmic tail) binding protein 2 (CD2BP2), mRNA
NM 006017	Homo sapiens prominin-like 1 (mouse) (PROML1), mRNA
NM_004010	Homo sapiens dystrophin (muscular dystrophy, Duchenne and Becker types), includes DXS142, DXS164, DXS206, DXS230, DXS239, DXS268, DXS269, DXS270, DXS272 (DMD), transcript variant Dp427p2, mRNA
NM_004023	Homo sapiens dystrophin (muscular dystrophy, Duchenne and Becker types), includes DXS142, DXS164, DXS206, DXS230, DXS239, DXS268, DXS269, DXS270, DXS272 (DMD), transcript variant Dp140bc, mRNA
NM_004022	Homo sapiens dystrophin (muscular dystrophy, Duchenne and Becker types), includes DXS142, DXS164, DXS206, DXS230, DXS239, DXS268, DXS269, DXS270, DXS272 (DMD), transcript variant D140ab, mRNA
NM_004021	Homo sapiens dystrophin (muscular dystrophy, Duchenne and Becker types), includes DXS142, DXS164, DXS206, DXS230, DXS239, DXS268, DXS269, DXS270, DXS272 (DMD), transcript variant Dp140b, mRNA

NM_004020	Homo sapiens dystrophin (muscular dystrophy, Duchenne and Becker types),
_	includes DXS142, DXS164, DXS206, DXS230, DXS239, DXS268, DXS269,
	DXS270, DXS272 (DMD), transcript variant Dp140c, mRNA
NM 004019	Homo sapiens dystrophin (muscular dystrophy, Duchenne and Becker types),
	includes DXS142, DXS164, DXS206, DXS230, DXS239, DXS268, DXS269,
	DXS270, DXS272 (DMD), transcript variant Dp40, mRNA
NM 004018	Homo sapiens dystrophin (muscular dystrophy, Duchenne and Becker types),
11171_004018	includes DXS142, DXS164, DXS206, DXS230, DXS239, DXS268, DXS269,
ND 6 004015	DXS270, DXS272 (DMD), transcript variant Dp71ab, mRNA
NM_004017	Homo sapiens dystrophin (muscular dystrophy, Duchenne and Becker types),
	includes DXS142, DXS164, DXS206, DXS230, DXS239, DXS268, DXS269,
	DXS270, DXS272 (DMD), transcript variant Dp71a, mRNA
NM_004016	Homo sapiens dystrophin (muscular dystrophy, Duchenne and Becker types),
	includes DXS142, DXS164, DXS206, DXS230, DXS239, DXS268, DXS269,
	DXS270, DXS272 (DMD), transcript variant Dp71b, mRNA
NM_004015	Homo sapiens dystrophin (muscular dystrophy, Duchenne and Becker types),
_	includes DXS142, DXS164, DXS206, DXS230, DXS239, DXS268, DXS269,
	DXS270, DXS272 (DMD), transcript variant Dp71, mRNA
NM 004014	Homo sapiens dystrophin (muscular dystrophy, Duchenne and Becker types),
00 101 7	includes DXS142, DXS164, DXS206, DXS230, DXS239, DXS268, DXS269,
	DXS270, DXS272 (DMD), transcript variant Dp116, mRNA
NM 004013	Homo sapiens dystrophin (muscular dystrophy, Duchenne and Becker types),
14141 ⁻ 004012	
	includes DXS142, DXS164, DXS206, DXS230, DXS239, DXS268, DXS269,
37.6.004040	DXS270, DXS272 (DMD), transcript variant Dp140, mRNA
NM_004012	Homo sapiens dystrophin (muscular dystrophy, Duchenne and Becker types),
	includes DXS142, DXS164, DXS206, DXS230, DXS239, DXS268, DXS269,
	DXS270, DXS272 (DMD), transcript variant Dp260-2, mRNA
NM_004011	Homo sapiens dystrophin (muscular dystrophy, Duchenne and Becker types),
	includes DXS142, DXS164, DXS206, DXS230, DXS239, DXS268, DXS269,
	DXS270, DXS272 (DMD), transcript variant Dp260-1, mRNA
NM_004009	Homo sapiens dystrophin (muscular dystrophy, Duchenne and Becker types),
_	includes DXS142, DXS164, DXS206, DXS230, DXS239, DXS268, DXS269,
	DXS270, DXS272 (DMD), transcript variant Dp427p1, mRNA
NM 004007	Homo sapiens dystrophin (muscular dystrophy, Duchenne and Becker types),
111.2_00 .00.	includes DXS142, DXS164, DXS206, DXS230, DXS239, DXS268, DXS269,
	DXS270, DXS272 (DMD), transcript variant Dp4271, mRNA
NM_004006	Homo sapiens dystrophin (muscular dystrophy, Duchenne and Becker types),
14141_00+000	includes DXS142, DXS164, DXS206, DXS230, DXS239, DXS268, DXS269,
NR (000100	DXS270, DXS272 (DMD), transcript variant Dp427m, mRNA
NM_000109	Homo sapiens dystrophin (muscular dystrophy, Duchenne and Becker types),
	includes DXS142, DXS164, DXS206, DXS230, DXS239, DXS268, DXS269,
	DXS270, DXS272 (DMD), transcript variant Dp427c, mRNA
NM_005657	Homo sapiens tumor protein p53 binding protein, 1 (TP53BP1), mRNA
NM_005632	Homo sapiens small optic lobes homolog (Drosophila) (SOLH), mRNA
NM_005631	Homo sapiens smoothened homolog (Drosophila) (SMOH), mRNA
NM 005621	Homo sapiens S100 calcíum binding protein A12 (calgranulin C) (S100A12),
	mRNA
NM 005620	Homo sapiens S100 calcium binding protein A11 (calgizzarin) (S100A11),
1111_005020	mRNA
NM 005610	
	Homo sapiens retinoblastoma binding protein 4 (RBBP4), mRNA
NM_005732	Homo sapiens RAD50 homolog (S. cerevisiae) (RAD50), mRNA
NM_005591	Homo sapiens MRE11 meiotic recombination 11 homolog A (S. cerevisiae)
	(MRE11A), mRNA

NM_005590	Homo sapiens MRE11 meiotic recombination 11 homolog A (S. cerevisiae) (MRE11A), mRNA
NM_005585	Homo sapiens MAD, mothers against decapentaplegic homolog 6 (Drosophila)
	(MADH6), mRNA
NM_005584	Homo sapiens mab-21-like 1 (C. elegans) (MAB21L1), mRNA
NM_005582	Homo sapiens lymphocyte antigen 64 homolog, radioprotective 105kD (mouse) (LY64), mRNA
NM 005667	Homo sapiens zinc finger protein 103 homolog (mouse) (ZFP103), mRNA
NM 005886	Homo sapiens katanin p80 (WD40-containing) subunit B 1 (KATNB1), mRNA
NM 005860	Homo sapiens follistatin-like 3 (secreted glycoprotein) (FSTL3), mRNA
NM 005758	Homo sapiens heterogeneous nuclear ribonucleoprotein A3 (HNRPA3), mRNA
NM_005510	Homo sapiens dom-3 homolog Z (C. elegans) (DOM3Z), transcript variant 2, mRNA
NM 005766	Homo sapiens FERM, RhoGEF (ARHGEF) and pleckstrin domain protein 1
	(chondrocyte-derived) (FARP1), mRNA
NM 005722	Homo sapiens ARP2 actin-related protein 2 homolog (yeast) (ACTR2), mRNA
NM 005750	Homo sapiens chromosome 4 open reading frame 6 (C4orf6), mRNA
NM_005170	Homo sapiens achaete-scute complex-like 2 (Drosophila) (ASCL2), mRNA
NM_005426	Homo sapiens tumor protein p53 binding protein, 2 (TP53BP2), mRNA
NM_005486	Homo sapiens target of myb1-like 1 (chicken) (TOM1L1), mRNA
NM_005488	Homo sapiens target of myb1 (chicken) (TOM1), mRNA
NM_005417	Homo sapiens v-src sarcoma (Schmidt-Ruppin A-2) viral oncogene homolog
_	(avian) (SRC), mRNA
NM_005413	Homo sapiens sine oculis homeobox homolog 3 (Drosophila) (SIX3), mRNA
NM_005444	Homo sapiens RCD1 required for cell differentiation1 homolog (S. pombe)
	(RQCD1), mRNA
NM_005378	Homo sapiens v-myc myelocytomatosis viral related oncogene, neuroblastoma derived (avian) (MYCN), mRNA
NM_005377	Homo sapiens v-myc myelocytomatosis viral oncogene homolog 2 (avian) (MYCL2), mRNA
NM_005375	Homo sapiens v-myb myeloblastosis viral oncogene homolog (avian) (MYB), mRNA
NM_005359	Homo sapiens MAD, mothers against decapentaplegic homolog 4 (Drosophila) (MADH4), mRNA
NM 005340	Homo sapiens histidine triad nucleotide binding protein (HINT), mRNA
NM_005307	Homo sapiens G protein-coupled receptor kinase 2-like (Drosophila) (GPRK2L), mRNA
NM_005262	Homo sapiens growth factor, augmenter of liver regeneration (ERV1 homolog, S. cerevisiae) (GFER), mRNA
NM_005261	Homo sapiens GTP binding protein overexpressed in skeletal muscle (GEM), mRNA
NM 005257	Homo sapiens GATA binding protein 6 (GATA6), mRNA
NM 005245	Homo sapiens FAT tumor suppressor homolog 1 (Drosophila) (FAT), mRNA
NM 005244	Homo sapiens eyes absent homolog 2 (Drosophila) (EYA2), mRNA
NM_005239	Homo sapiens v-ets erythroblastosis virus E26 oncogene homolog 2 (avian) (ETS2), mRNA
NM_005235	Homo sapiens v-erb-a erythroblastic leukemia viral oncogene homolog 4 (avian) (ERBB4), mRNA
NM_005228	Homo sapiens epidermal growth factor receptor (erythroblastic leukemia viral (verb-b) oncogene homolog, avian) (EGFR), mRNA
NM 005224	Homo sapiens dead ringer-like 1 (Drosophila) (DRIL1), mRNA
NM 005219	Homo sapiens dead iniger-like I (Diosophila) (DIAPHI), mRNA
14141 003213	Trome suprems diaphianous nomiolog I (Drosophina) (Dia III), mad it

NM_005207	Homo sapiens v-crk sarcoma virus CT10 oncogene homolog (avian)-like (CRKL), mRNA
NM_005197	Homo sapiens checkpoint suppressor 1 (CHES1), mRNA
NM 005454	Homo sapiens cerberus 1 homolog, cysteine knot superfamily (Xenopus laevis)
	(CER1), mRNA
NM 005496	Homo sapiens SMC4 structural maintenance of chromosomes 4-like 1 (yeast)
_	(SMC4L1), mRNA
NM 005169	Homo sapiens aristaless homeobox (Drosophila) (ARIX), mRNA
NM_005078	Homo sapiens transducin-like enhancer of split 3 (E(sp1) homolog, Drosophila)
	(TLE3), mRNA
NM_005077	Homo sapiens transducin-like enhancer of split 1 (E(sp1) homolog, Drosophila)
	(TLE1), mRNA
NM_005068	Homo sapiens single-minded homolog 1 (Drosophila) (SIM1), mRNA
NM_005067	Homo sapiens seven in absentia homolog 2 (Drosophila) (SIAH2), mRNA
NM_005138	Homo sapiens SCO cytochrome oxidase deficient homolog 2 (yeast) (SCO2),
	nuclear gene encoding mitochondrial protein, mRNA
NM_005156	Homo sapiens ROD1 regulator of differentiation 1 (S. pombe) (ROD1), mRNA
NM_005133	Homo sapiens RCE1 homolog, prenyl protein protease (S. cerevisiae) (RCE1),
	mRNA DE LA CORRECCIONA DEL CORRECCIONA DE LA CORRECCIONA DEL CORRECCIONA DE LA CORRECCIONA DEL CORRECCIONA DE LA CORRECCIONA DEL CORRECCIONA DE LA CORRECCIO
NM_005057	Homo sapiens retinoblastoma binding protein 5 (RBBP5), mRNA
NM_005056	Homo sapiens retinoblastoma binding protein 2 (RBBP2), mRNA
NM_005053	Homo sapiens RAD23 homolog A (S. cerevisiae) (RAD23A), mRNA
NM_005049	Homo sapiens PWP2 periodic tryptophan protein homolog (yeast) (PWP2H), mRNA
NM_005008	Homo sapiens NHP2 non-histone chromosome protein 2-like 1 (S. cerevisiae)
	(NHP2L1), mRNA
NM_004997	Homo sapiens myosin binding protein H (MYBPH), mRNA
NM_004677	Homo sapiens Testis-specific XK-related protein on Y (XKRY), mRNA
NM_004788	Homo sapiens ubiquitination factor E4A (UFD2 homolog, yeast) (UBE4A), mRNA
NM_004617	Homo sapiens transmembrane 4 superfamily member 4 (TM4SF4), mRNA
NM_004607	Homo sapiens tubulin-specific chaperone a (TBCA), mRNA
NM_004602	Homo sapiens staufen, RNA binding protein (Drosophila) (STAU), transcript variant T4, mRNA
NM_004653	Homo sapiens Smcy homolog, Y chromosome (mouse) (SMCY), mRNA
NM_004787	Homo sapiens slit homolog 2 (Drosophila) (SLIT2), mRNA
NM_004593	Homo sapiens splicing factor, arginine/serine-rich 10 (transformer 2 homolog, Drosophila) (SFRS10), mRNA
NM 004206	Homo sapiens vesicle trafficking protein (SEC22C), transcript variant 2, mRNA
NM 004657	Homo sapiens serum deprivation response (phosphatidylserine binding protein)
_	(SDPR), mRNA
NM_004589	Homo sapiens SCO cytochrome oxidase deficient homolog 1 (yeast) (SCO1),
	nuclear gene encoding mitochondrial protein, mRNA
NM_004587	Homo sapiens ribosome binding protein 1 homolog 180kD (dog) (RRBP1),
	mRNA
NM_004164	Homo sapiens retinol binding protein 2, cellular (RBP2), mRNA
NM_004584	Homo sapiens RAD9 homolog (S. pombe) (RAD9), mRNA
NM_004794	Homo sapiens RAB33A, member RAS oncogene family (RAB33A), mRNA
NM_004813	Homo sapiens peroxisomal biogenesis factor 16 (PEX16), transcript variant 1, mRNA
NM 004564	Homo sapiens PET112-like (yeast) (PET112L), mRNA
NM 004643	Homo sapiens PETTIZ-like (yeast) (PETTIZE), interval Homo sapiens poly(A) binding protein, nuclear 1 (PABPN1), mRNA
14141 004043	1 Homo sapiens pory(A) uniquing protein, nuclear 1 (1702247), find A

NDC 0045C1	Homo sapiens ovo-like 1(Drosophila) (OVOL1), mRNA
NM_004561	Homo sapiens ovo-like (Diosophila) (OVOLI), likeva Homo sapiens origin recognition complex, subunit 1-like (yeast) (ORC1L),
NM_004153	mRNA
NM 004557	Homo sapiens Notch homolog 4 (Drosophila) (NOTCH4), mRNA
NM 004808	Homo sapiens N-myristoyltransferase 2 (NMT2), mRNA
NM 004210	Homo sapiens neuralized-like (Drosophila) (NEURL), mRNA
NM 004147	Homo sapiens developmentally regulated GTP binding protein 1 (DRG1),
	mRNA
NM_004851	Homo sapiens pronapsin A (NAP1), mRNA
NM_004533	Homo sapiens myosin binding protein C, fast type (MYBPC2), mRNA
NM_004529	Homo sapiens myeloid/lymphoid or mixed-lineage leukemia (trithorax homolog,
	Drosophila); translocated to, 3 (MLLT3), mRNA
NM_004668	Homo sapiens maltase-glucoamylase (alpha-glucosidase) (MGAM), mRNA
NM_004526	Homo sapiens MCM2 minichromosome maintenance deficient 2, mitotin (S. cerevisiae) (MCM2), mRNA
NM_004829	Homo sapiens lymphocyte antigen 94 homolog, activating NK-receptor; NK-
	p46, (mouse) (LY94), mRNA
NM_004744	Homo sapiens lecithin retinol acyltransferase (phosphatidylcholineretinol O-
	acyltransferase) (LRAT), mRNA
NM_004524	Homo sapiens lethal giant larvae homolog 2 (Drosophila) (LLGL2), mRNA
NM_004140	Homo sapiens lethal giant larvae homolog 1 (Drosophila) (LLGL1), mRNA
NM_004922	Homo sapiens SEC24 related gene family, member C (S. cerevisiae) (SEC24C), mRNA
NM 004508	Homo sapiens isopentenyl-diphosphate delta isomerase (IDI1), mRNA
NM 004507	Homo sapiens HUS1 checkpoint homolog (S. pombe) (HUS1), mRNA
NM 004262	Homo sapiens airway trypsin-like protease (HAT), mRNA
NM 004752	Homo sapiens glial cells missing homolog b (Drosophila) (GCMB), mRNA
NM 004477	Homo sapiens FSHD region gene 1 (FRG1), mRNA
NM 004463	Homo sapiens faciogenital dysplasia (Aarskog-Scott syndrome) (FGD1), mRNA
NM 004106	Homo sapiens Fc fragment of IgE, high affinity I, receptor for; gamma
1411_00 1100	polypeptide (FCER1G), mRNA
NM_004456	Homo sapiens enhancer of zeste homolog 2 (Drosophila) (EZH2), mRNA
NM_004100	Homo sapiens eyes absent homolog 4 (Drosophila) (EYA4), mRNA
NM 004450	Homo sapiens enhancer of rudimentary homolog (Drosophila) (ERH), mRNA
NM_004448	Homo sapiens v-erb-b2 erythroblastic leukemia viral oncogene homolog 2,
	neuro/glioblastoma derived oncogene homolog (avian) (ERBB2), mRNA
NM_004445	Homo sapiens EphB6 (EPHB6), mRNA
NM_004436	Homo sapiens endosulfine alpha (ENSA), mRNA
NM_004432	Homo sapiens ELAV (embryonic lethal, abnormal vision, Drosophila)-like 2 (Hu
	antigen B) (ELAVL2), mRNA
NM_004230	Homo sapiens endothelial differentiation, sphingolipid G-protein-coupled
	receptor, 5 (EDG5), mRNA
NM_004421	Homo sapiens dishevelled, dsh homolog 1 (Drosophila) (DVL1), mRNA
NM_004399	Homo sapiens DEAD/H (Asp-Glu-Ala-Asp/His) box polypeptide 11 (CHL1-like
	helicase homolog, S. cerevisiae) (DDX11), transcript variant 2, mRNA
NM_004378	Homo sapiens cellular retinoic acid binding protein 1 (CRABP1), mRNA
NM_004898	Homo sapiens clock homolog (mouse) (CLOCK), mRNA
NM_004669	Homo sapiens chloride intracellular channel 3 (CLIC3), mRNA
NM_004066	Homo sapiens centrin, EF-hand protein, 1 (CETN1), mRNA
NM_004354	Homo sapiens cyclin G2 (CCNG2), mRNA
NM_004352	Homo sapiens cerebellin 1 precursor (CBLN1), mRNA
NM 004057	Homo sapiens calbindin 3, (vitamin D-dependent calcium binding protein)

	(CALB3), mRNA
	Homo sapiens chromosome 18 open reading frame 1 (C18orf1), mRNA
NM_004725 I	Homo sapiens BUB3 budding uninhibited by benzimidazoles 3 homolog (yeast)
	(BUB3), mRNA
NM 004336	Homo sapiens BUB1 budding uninhibited by benzimidazoles 1 homolog (yeast)
	(BUB1), mRNA
	Homo sapiens BCL2/adenovirus E1B 19kD interacting protein 3-like (BNIP3L),
	mRNA
	Homo sapiens BCS1-like (yeast) (BCS1L), mRNA
NM 004045 I	Homo sapiens ATX1 antioxidant protein 1 homolog (yeast) (ATOX1), mRNA
NM 004849 I	Homo sapiens APG5 autophagy 5-like (S. cerevisiae) (APG5L), mRNA
NM_004674 I	Homo sapiens ash2 (absent, small, or homeotic)-like (Drosophila) (ASH2L),
	mRNA
	Homo sapiens achaete-scute complex-like 1 (Drosophila) (ASCL1), mRNA
	Homo sapiens APG12 autophagy 12-like (S. cerevisiae) (APG12L), mRNA
	Homo sapiens myeloid/lymphoid or mixed-lineage leukemia (trithorax homolog,
	Drosophila); translocated to, 10 (MLLT10), mRNA
	Homo sapiens BAF53 (BAF53A), mRNA
	Homo sapiens AE binding protein 1 (AEBP1), mRNA
	Homo sapiens calcium/calmodulin-dependent protein kinase I (CAMK1), mRNA
	Homo sapiens lysozyme (renal amyloidosis) (LYZ), mRNA
	Homo sapiens sulfite oxidase (SUOX), nuclear gene encoding mitochondrial
	protein, mRNA
	Homo sapiens Notch homolog 3 (Drosophila) (NOTCH3), mRNA
	Homo sapiens mutS homolog 2, colon cancer, nonpolyposis type 1 (E. coli) (MSH2), mRNA
	Homo sapiens mutL homolog 1, colon cancer, nonpolyposis type 2 (E. coli)
	(MLH1), mRNA
NM_000210	Homo sapiens integrin, alpha 6 (ITGA6), mRNA
NM_001537	Homo sapiens heat shock factor binding protein 1 (HSBP1), mRNA
NM_001499	Homo sapiens GLE1 RNA export mediator-like (yeast) (GLE1L), mRNA
	Homo sapiens filamin C, gamma (actin binding protein 280) (FLNC), mRNA
	Homo sapiens fatty acid binding protein 5 (psoriasis-associated) (FABP5), mRNA
	Homo sapiens epiregulin (EREG), mRNA
NM_001388	Homo sapiens developmentally regulated GTP binding protein 2 (DRG2), mRNA
	Homo sapiens cylicin, basic protein of sperm head cytoskeleton 2 (CYLC2),
	mRNA
	Homo sapiens cleavage stimulation factor, 3' pre-RNA, subunit 3, 77kD
_	(CSTF3), mRNA
	Homo sapiens cleavage stimulation factor, 3' pre-RNA, subunit 2, 64kD
	(CSTF2), mRNA
	Homo sapiens cleavage stimulation factor, 3' pre-RNA, subunit 1, 50kD
1 - 1	(CSTF1), mRNA
	Homo sapiens CDC20 cell division cycle 20 homolog (S. cerevisiae) (CDC20),
	mRNA
NM_001122	Homo sapiens adipose differentiation-related protein (ADFP), mRNA
NM_003413	Homo sapiens Zic family member 3 heterotaxy 1 (odd-paired homolog,
	Drosophila) (ZIC3), mRNA
NM_003412	Homo sapiens Zic family member 1 (odd-paired homolog, Drosophila) (ZIC1),
	mRNA

NM_003408	Homo sapiens zinc finger protein 37 homolog (mouse) (ZFP37), mRNA
NM_003409	Homo sapiens zinc finger protein 161 homolog (mouse) (ZFP161), mRNA
NM_003680	Homo sapiens tyrosyl-tRNA synthetase (YARS), mRNA
NM_003390	Homo sapiens WEE1+ homolog (S. pombe) (WEE1), mRNA
NM_003565	Homo sapiens unc-51-like kinase 1 (C. elegans) (ULK1), mRNA
NM_003345	Homo sapiens ubiquitin-conjugating enzyme E2I (UBC9 homolog, yeast)
	(UBE2I), mRNA
NM_003344	Homo sapiens ubiquitin-conjugating enzyme E2H (UBC8 homolog, yeast)
275 0000 10	(UBE2H), mRNA
NM_003343	Homo sapiens ubiquitin-conjugating enzyme E2G 2 (UBC7 homolog, yeast)
37 C 0000 40	(UBE2G2), mRNA
NM_003340	Homo sapiens ubiquitin-conjugating enzyme E2D 3 (UBC4/5 homolog, yeast)
ND4 002220	(UBE2D3), mRNA
NM_003338	Homo sapiens ubiquitin-conjugating enzyme E2D 1 (UBC4/5 homolog, yeast) (UBE2D1), mRNA
NM_003968	Homo sapiens ubiquitin-activating enzyme E1C (UBA3 homolog, yeast)
NM 003320	(UBE1C), mRNA
NM_003278	Homo sapiens tubby homolog (mouse) (TUB), mRNA
NM 003278	Homo sapiens tetranectin (plasminogen binding protein) (TNA), mRNA
14M_003200	Homo sapiens transducin-like enhancer of split 2 (E(sp1) homolog, Drosophila) (TLE2), mRNA
NM 003920	
NM 003251	Homo sapiens timeless homolog (Drosophila) (TIMELESS), mRNA
	Homo sapiens thyroid hormone responsive (SPOT14 homolog, rat) (THRSP), mRNA
NM_003250	Homo sapiens thyroid hormone receptor, alpha (erythroblastic leukemia viral (v-
	erb-a) oncogene homolog, avian) (THRA), mRNA
NM_003223	Homo sapiens transcription factor AP-4 (activating enhancer binding protein 4)
	(TFAP4), mRNA
NM_003222	Homo sapiens transcription factor AP-2 gamma (activating enhancer binding
	protein 2 gamma) (TFAP2C), mRNA
NM_003221	Homo sapiens transcription factor AP-2 beta (activating enhancer binding protein
	2 beta) (TFAP2B), mRNA
NM_003220	Homo sapiens transcription factor AP-2 alpha (activating enhancer binding
277 600450	protein 2 alpha) (TFAP2A), mRNA
NM_000458	Homo sapiens transcription factor 2, hepatic; LF-B3; variant hepatic nuclear
ND4 002101	factor (TCF2), transcript variant a, mRNA
NM_003181 NM_003173	Homo sapiens T, brachyury homolog (mouse) (T), mRNA
1N1VI_003173	Homo sapiens suppressor of variegation 3-9 homolog 1 (Drosophila)
NM 003171	(SUV39H1), mRNA
NM 003169	Homo sapiens suppressor of var1, 3-like 1 (S. cerevisiae) (SUPV3L1), mRNA
NM_003168	Homo sapiens suppressor of Ty 5 homolog (S. cerevisiae) (SUPT5H), mRNA
NM 003599	Homo sapiens suppressor of Ty 4 homolog 1 (S. cerevisiae) (SUPT4H1), mRNA
NM_003162	Homo sapiens suppressor of Ty 3 homolog (S. cerevisiae) (SUPT3H), mRNA
NM_003134	Homo sapiens striatin, calmodulin binding protein (STRN), mRNA
	Homo sapiens signal recognition particle 14kD (homologous Alu RNA binding protein) (SRP14), mRNA
NM_003088	Homo sapiens singed-like (fascin homolog, sea urchin) (Drosophila) (SNL),
	mRNA
NM_003061	Homo sapiens slit homolog 1 (Drosophila) (SLIT1), mRNA
NM_003036	Homo sapiens v-ski sarcoma viral oncogene homolog (avian) (SKI), mRNA
NM_003031	Homo sapiens seven in absentia homolog 1 (Drosophila) (SIAH1), mRNA
NM_000193	Homo sapiens sonic hedgehog homolog (Drosophila) (SHH), mRNA
	woderiog nomotog (Diosophina) (Sitti), liikiva

NM_003003	Homo sapiens SEC14-like 1 (S. cerevisiae) (SEC14L1), mRNA
NM_002983	Homo sapiens small inducible cytokine A3 (SCYA3), mRNA
NM_002982	Homo sapiens small inducible cytokine A2 (monocyte chemotactic protein 1) (SCYA2), mRNA
NM 002981	Homo sapiens small inducible cytokine A1, I-309 (SCYA1), mRNA
NM 003864	Homo sapiens sin3-associated polypeptide, 30kD (SAP30), mRNA
NM 002962	Homo sapiens S100 calcium binding protein A5 (S100A5), mRNA
NM 002960	Homo sapiens S100 calcium binding protein A3 (S100A3), mRNA
NM 002966	Homo sapiens S100 calcium binding protein A10 (annexin II ligand, calpactin I,
	light polypeptide (p11)) (S100A10), mRNA
NM_003707	Homo sapiens RuvB-like 1 (E. coli) (RUVBL1), mRNA
NM_002944	Homo sapiens v-ros UR2 sarcoma virus oncogene homolog 1 (avian) (ROS1), mRNA
NM_002941	Homo sapiens roundabout, axon guidance receptor, homolog 1 (Drosophila) (ROBO1), mRNA
NM_000326	Homo sapiens retinaldehyde binding protein 1 (RLBP1), mRNA
NM 002930	Homo sapiens Ric-like, expressed in neurons (Drosophila) (RIN), mRNA
NM_003961	Homo sapiens rhomboid, veinlet-like 1 (Drosophila) (RHBDL), mRNA
NM_002912	Homo sapiens REV3-like, catalytic subunit of DNA polymerase zeta (yeast) (REV3L), mRNA
NM 002900	Homo sapiens retinol binding protein 3, interstitial (RBP3), mRNA
NM 002894	Homo sapiens retinoblastoma binding protein 8 (RBBP8), mRNA
NM_002888	Homo sapiens retinoic acid receptor responder (tazarotene induced) 1 (RARRES1), mRNA
NM 002879	Homo sapiens RAD52 homolog (S. cerevisiae) (RAD52), mRNA
NM 002878	Homo sapiens RAD51-like 3 (S. cerevisiae) (RAD51L3), mRNA
NM_002875	Homo sapiens RAD51 homolog (RecA homolog, E. coli) (S. cerevisiae) (RAD51), mRNA
NM 002874	Homo sapiens RAD23 homolog B (S. cerevisiae) (RAD23B), mRNA
NM 002853	Homo sapiens RAD1 homolog (S. pombe) (RAD1), mRNA
NM 002873	Homo sapiens RAD17 homolog (S. pombe) (RAD17), mRNA
NM 000264	Homo sapiens patched homolog (Drosophila) (PTCH), mRNA
NM 003738	Homo sapiens patched homolog 2 (Drosophila) (PTCH2), mRNA
NM 002616	Homo sapiens period homolog 1 (Drosophila) (PER1), mRNA
NM_002600	Homo sapiens phosphodiesterase 4B, cAMP-specific (phosphodiesterase E4
	dunce homolog, Drosophila) (PDE4B), mRNA
NM 002568	Homo sapiens poly(A) binding protein, cytoplasmic 1 (PABPC1), mRNA
NM_003932	Homo sapiens suppression of tumorigenicity 13 (colon carcinoma) (Hsp70 interacting protein) (ST13), mRNA
NM 003715	Homo sapiens vesicle docking protein p115 (P115), mRNA
NM_002553	Homo sapiens origin recognition complex, subunit 5-like (yeast) (ORC5L), mRNA
NM_002552	Homo sapiens origin recognition complex, subunit 4-like (yeast) (ORC4L), mRNA
NM 003634	Homo sapiens nipsnap homolog 1 (C. elegans) (NIPSNAP1), mRNA
NM 002499	Homo sapiens neogenin homolog 1 (chicken) (NEO1), mRNA
NM_002484	Homo sapiens nucleotide binding protein 1 (MinD homolog, E. coli) (NUBP1), mRNA
NM_003827	Homo sapiens N-ethylmaleimide-sensitive factor attachment protein, alpha (NAPA), mRNA
NM_002466	Homo sapiens v-myb myeloblastosis viral oncogene homolog (avian)-like 2 (MYBL2), mRNA

Iomo sapiens msh homeo box homolog 1 (Drosophila) (MSX1), mRNA
Iomo sapiens serine/threonine kinase 24 (STE20 homolog, yeast) (STK24),
nRNA
Iomo sapiens musashi homolog 1 (Drosophila) (MSI1), mRNA
Iomo sapiens mutS homolog 5 (E. coli) (MSH5), mRNA
Iomo sapiens mutS homolog 4 (E. coli) (MSH4), mRNA
Iomo sapiens mutS homolog 3 (E. coli) (MSH3), mRNA
Iomo sapiens manic fringe homolog (Drosophila) (MFNG), mRNA
Iomo sapiens mesoderm specific transcript homolog (mouse) (MEST), mRNA
Iomo sapiens Meis1, myeloid ecotropic viral integration site 1 homolog (mouse) MEIS1), mRNA
Homo sapiens Mdm4, transformed 3T3 cell double minute 4, p53 binding protein mouse) (MDM4), mRNA
Homo sapiens Mdm2, transformed 3T3 cell double minute 2, p53 binding protein mouse) (MDM2), transcript variant MDM2, mRNA
Homo sapiens MCM3 minichromosome maintenance deficient 3 (S. cerevisiae) ssociated protein (MCM3AP), mRNA
Homo sapiens v-maf musculoaponeurotic fibrosarcoma oncogene homolog Kavian) (MAFK), mRNA
Homo sapiens v-maf musculoaponeurotic fibrosarcoma oncogene homolog Gavian) (MAFG), mRNA
Homo sapiens MAD1 mitotic arrest deficient-like 1 (yeast) (MAD1L1), mRNA
Iomo sapiens kynureninase (L-kynurenine hydrolase) (KYNU), mRNA
Homo sapiens karyopherin alpha 5 (importin alpha 6) (KPNA5), mRNA
Homo sapiens jerky homolog-like (mouse) (JRKL), mRNA
Homo sapiens ISL1 transcription factor, LIM/homeodomain, (islet-1) (ISL1), nRNA
Iomo sapiens insulin receptor substrate 4 (IRS4), mRNA
Iomo sapiens interleukin-1 receptor-associated kinase 2 (IRAK2), mRNA
Homo sapiens inositol polyphosphate-4-phosphatase, type II, 105kD (INPP4B), nRNA
Homo sapiens HMT1 hnRNP methyltransferase-like 2 (S. cerevisiae) HRMT1L2), mRNA
Homo sapiens HMT1 hnRNP methyltransferase-like 1 (S. cerevisiae) HRMT1L1), mRNA
Homo sapiens harakiri, BCL2 interacting protein (contains only BH3 domain) HRK), mRNA
Homo sapiens histidine rich calcium binding protein (HRC), mRNA
Homo sapiens human immunodeficiency virus type I enhancer binding protein 1 HIVEP1), mRNA
Homo sapiens serine protease inhibitor, Kunitz type 1 (SPINT1), mRNA
forms saprens serme procease minorior, Kumez type 1 (SI HVII), mktvA
Homo sapiens mutS homolog 6 (E. coli) (MSH6), mRNA
Homo sapiens mutS homolog 6 (E. coli) (MSH6), mRNA Homo sapiens glutamate receptor, metabotropic 2 (GRM2), mRNA
Homo sapiens mutS homolog 6 (E. coli) (MSH6), mRNA Homo sapiens glutamate receptor, metabotropic 2 (GRM2), mRNA Homo sapiens golgi autoantigen, golgin subfamily a, 1 (GOLGA1), mRNA
Homo sapiens mutS homolog 6 (E. coli) (MSH6), mRNA Homo sapiens glutamate receptor, metabotropic 2 (GRM2), mRNA Homo sapiens golgi autoantigen, golgin subfamily a, 1 (GOLGA1), mRNA Homo sapiens gamma-glutamyl hydrolase (conjugase, folylpolygammaglutamyl nydrolase) (GGH), mRNA
Homo sapiens mutS homolog 6 (E. coli) (MSH6), mRNA Homo sapiens glutamate receptor, metabotropic 2 (GRM2), mRNA Homo sapiens golgi autoantigen, golgin subfamily a, 1 (GOLGA1), mRNA Homo sapiens gamma-glutamyl hydrolase (conjugase, folylpolygammaglutamyl nydrolase) (GGH), mRNA Homo sapiens transcriptional adaptor 2 (ADA2 homolog, yeast)-like (TADA2L), mRNA
Homo sapiens mutS homolog 6 (E. coli) (MSH6), mRNA Homo sapiens glutamate receptor, metabotropic 2 (GRM2), mRNA Homo sapiens golgi autoantigen, golgin subfamily a, 1 (GOLGA1), mRNA Homo sapiens gamma-glutamyl hydrolase (conjugase, folylpolygammaglutamyl nydrolase) (GGH), mRNA Homo sapiens transcriptional adaptor 2 (ADA2 homolog, yeast)-like (TADA2L),
Homo sapiens mutS homolog 6 (E. coli) (MSH6), mRNA Homo sapiens glutamate receptor, metabotropic 2 (GRM2), mRNA Homo sapiens golgi autoantigen, golgin subfamily a, 1 (GOLGA1), mRNA Homo sapiens gamma-glutamyl hydrolase (conjugase, folylpolygammaglutamyl hydrolase) (GGH), mRNA Homo sapiens transcriptional adaptor 2 (ADA2 homolog, yeast)-like (TADA2L), mRNA Homo sapiens GCN5 general control of amino-acid synthesis 5-like 1 (yeast)

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NM_002051	Homo sapiens GATA binding protein 3 (GATA3), mRNA
NM_002050	Homo sapiens GATA binding protein 2 (GATA2), mRNA
NM_002049	Homo sapiens GATA binding protein 1 (globin transcription factor 1) (GATA1), mRNA
NM 002040	Homo sapiens GA binding protein transcription factor, alpha subunit (60kD)
	(GABPA), mRNA
NM_002039	Homo sapiens GRB2-associated binding protein 1 (GAB1), mRNA
NM_003508	Homo sapiens frizzled homolog 9 (Drosophila) (FZD9), mRNA
NM_003507	Homo sapiens frizzled homolog 7 (Drosophila) (FZD7), mRNA
NM_003506	Homo sapiens frizzled homolog 6 (Drosophila) (FZD6), mRNA
NM_003468	Homo sapiens frizzled homolog 5 (Drosophila) (FZD5), mRNA
NM_003505	Homo sapiens frizzled homolog 1 (Drosophila) (FZD1), mRNA
NM_001465	Homo sapiens FYN binding protein (FYB-120/130) (FYB), mRNA
NM_002031	Homo sapiens fyn-related kinase (FRK), mRNA
NM_003717	Homo sapiens neuropeptide FF-amide peptide precursor (NPFF), mRNA
NM_001457	Homo sapiens filamin B, beta (actin binding protein 278) (FLNB), mRNA
NM_001456	Homo sapiens filamin A, alpha (actin binding protein 280) (FLNA), mRNA
NM_002018	Homo sapiens flightless I homolog (Drosophila) (FLII), mRNA
NM_001991	Homo sapiens enhancer of zeste homolog 1 (Drosophila) (EZH1), mRNA
NM_001990	Homo sapiens eyes absent homolog 3 (Drosophila) (EYA3), mRNA
NM_000503	Homo sapiens eyes absent homolog 1 (Drosophila) (EYA1), mRNA
NM_001989	Homo sapiens eve, even-skipped homeo box homolog 1 (Drosophila) (EVX1), mRNA
NM 001982	Homo sapiens v-erb-b2 erythroblastic leukemia viral oncogene homolog 3
	(avian) (ERBB3), mRNA
NM_003584	Homo sapiens dual specificity phosphatase 11 (RNA/RNP complex 1-
	interacting) (DUSP11), mRNA
NM_003859	Homo sapiens dolichyl-phosphate mannosyltransferase polypeptide 1, catalytic subunit (DPM1), mRNA
NM 001928	Homo sapiens D component of complement (adipsin) (DF), mRNA
NM 003649	Homo sapiens D-aspartate oxidase (DDO), transcript variant 1, mRNA
NM_001343	Homo sapiens disabled homolog 2, mitogen-responsive phosphoprotein (Drosophila) (DAB2), mRNA
NM_001913	Homo sapiens cut-like 1, CCAAT displacement protein (Drosophila) (CUTL1), mRNA
NM 001316	Homo sapiens CSE1 chromosome segregation 1-like (yeast) (CSE1L), mRNA
NM 003652	Homo sapiens carboxypeptidase Z (CPZ), mRNA
NM 003909	Homo sapiens copine III (CPNE3), mRNA
NM 003915	Homo sapiens copine II (CPNE1), mRNA
NM_001308	Homo sapiens copine i (CINEI), indexA Homo sapiens carboxypeptidase N, polypeptide 1, 50kD (CPN1), mRNA
NM 001841	Homo sapiens cannabinoid receptor 2 (macrophage) (CNR2), mRNA
NM_001280	Homo sapiens cold inducible RNA binding protein (CIRBP), mRNA
NM_001274	Homo sapiens CHK1 checkpoint homolog (S. pombe) (CHEK1), mRNA
NM_001806	Homo sapiens CCAAT/enhancer binding protein (C/EBP), gamma (CEBPG),
	mRNA
NM_003655	Homo sapiens chromobox homolog 4 (Pc class homolog, Drosophila) (CBX4), mRNA
NM_001749	
NM 000716	Homo sapiens calpain, small subunit 1 (CAPNS1), mRNA Homo sapiens complement component 4 binding protein, beta (C4BPB), mRNA
	Homo sapiens complement component 4 binding protein, beta (C4BPA), likNA Homo sapiens complement component 4 binding protein, alpha (C4BPA),
NM_000715	mRNA
NM_001726	Homo sapiens bromodomain, testis-specific (BRDT), mRNA

ND 6 001005	Homo sapiens BCL2/adenovirus E1B 19kD interacting protein 1 (BNIP1),
NM_001205	transcript variant BNIP1, mRNA
NM 001714	Homo sapiens Bicaudal D homolog 1 (Drosophila) (BICD1), mRNA
NM 003766	Homo sapiens beclin 1 (coiled-coil, myosin-like BCL2 interacting protein)
	(BECN1), mRNA
NM 003567	Homo sapiens breast cancer anti-estrogen resistance 3 (BCAR3), mRNA
NM_001189	Homo sapiens bagpipe homeobox homolog 1 (Drosophila) (BAPX1), mRNA
NM 001698	Homo sapiens AU RNA binding protein/enoyl-Coenzyme A hydratase (AUH),
112/2_002050	nuclear gene encoding mitochondrial protein, mRNA
NM 001672	Homo sapiens agouti signaling protein, nonagouti homolog (mouse) (ASIP),
	mRNA
NM 001638	Homo sapiens apolipoprotein F (APOF), mRNA
NM 003977	Homo sapiens aryl hydrocarbon receptor interacting protein (AIP), mRNA
NM 001138	Homo sapiens agouti related protein homolog (mouse) (AGRP), transcript
	variant 1, mRNA
NM 058246	Homo sapiens DnaJ (Hsp40) homolog, subfamily B, member 6 (DNAJB6),
_	mRNA
NM 025225	Homo sapiens hypothetical protein dJ796I17.1 (DJ796I17.1), mRNA
NM 058165	Homo sapiens diacylglycerol acyltransferase 2-like (DGAT2-like), mRNA
NM 001861	Homo sapiens cytochrome c oxidase subunit IV isoform 1 (COX4I1), nuclear
_	gene encoding mitochondrial protein, mRNA
NM 014491	Homo sapiens forkhead box P2 (FOXP2), mRNA
NM 054110	Homo sapiens UDP-N-acetyl-alpha-D-galactosamine:polypeptide N-
	acetylgalactosaminyltransferase 7 (GALNT7), mRNA
NM 006726	Homo sapiens vesicle trafficking, beach and anchor containing (LRBA), mRNA
NM 020663	Homo sapiens TC10-like Rho GTPase (TCL), mRNA
NM 020919	Homo sapiens amyotrophic lateral sclerosis 2 (juvenile) (ALS2), mRNA
NM 052852	Homo sapiens hypothetical zinc finger protein MGC2396 (MGC2396), mRNA
NM 053043	Homo sapiens hypothetical protein MGC20460 (MGC20460), mRNA
NM 053017	Homo sapiens ADP-ribosyltransferase 5 (ART5), mRNA
NM 052999	Homo sapiens chemokine-like factor-like protein CKLFH1 (CKLFH1), mRNA
NM_052881	Homo sapiens hypothetical protein dJ734P14.5 (novel C2H2 type zinc finger
	protein) (MGC20504), mRNA
NM_052968	Homo sapiens apolipoprotein A-V (APOA5), mRNA
NM_052960	Homo sapiens retinoid binding protein 7 (RBP7), mRNA
NM_052959	Homo sapiens pannexin 3 (PANX3), mRNA
NM_052948	Homo sapiens sorting nexin 26 (SNX26), mRNA
NM_052947	Homo sapiens heart alpha-kinase (HAK), mRNA
NM_052946	Homo sapiens hypothetical protein MGC20702 (MGC20702), mRNA
NM_052943	Homo sapiens hypothetical protein MGC16491 (MGC16491), mRNA
NM_052941	Homo sapiens guanylate binding protein 4 (GBP4), mRNA
NM_052935	Homo sapiens hypothetical protein MGC20781 (MGC20781), mRNA
NM_052890	Homo sapiens peptidoglycan recognition protein L precursor (PGLYRP), mRNA
NM_052885	Homo sapiens solute carrier family 2 (facilitated glucose transporter), member 13
	(SLC2A13), mRNA
NM_052884	Homo sapiens sialic acid binding Ig-like lectin 11 (SIGLEC11), mRNA
NM_052877	Homo sapiens similar to hypothetical protein MNCb-2386 (MGC17544), mRNA
NM_052876	Homo sapiens transcriptional repressor NAC1 (NAC1), mRNA
NM_052873	Homo sapiens MGC16028 similar to RIKEN cDNA 1700019E19 gene
	(MGC16028), mRNA
NM_052871	Homo sapiens hypothetical protein MGC4677 (MGC4677), mRNA
NM_052870	Homo sapiens sorting nexin 18 (SNX18), mRNA

324 050050	
NM_052859	Homo sapiens putative endoplasmic reticulum multispan transmembrane protein
NM 052858	(RFT1), mRNA Homo sapiens similar to RIKEN cDNA 1810006A16 gene (LOC91862), mRNA
NM 052855	Homo sapiens hypothetical protein MGC15396 (MGC15396), mRNA
NM 052854	Homo sapiens old astrocyte specifically induced substance (OASIS), mRNA
NM 052844	Homo sapiens hypothetical protein MGC20486 (MGC20486), mRNA
NM 052839	Homo sapiens pannexin 2 (PANX2), mRNA
NM 033551	Homo sapiens hypothetical protein MGC19556 (MGC19556), mRNA
NM 033549	Homo sapiens hypothetical gene MGC1127 (MGC13530), mRNA
NM 033546	Homo sapiens myosin regulatory light chain (MLC-B), mRNA
NM 033544	Homo sapiens similar to cyclin-E binding protein 1 (H. sapiens) (MGC14386),
	mRNA
NM 033515	Homo sapiens MacGAP protein (MacGAP), mRNA
NM 033519	Homo sapiens olfactory receptor sdolf (sdolf), mRNA
NM 033516	Homo sapiens protein kinase NYD-SP25 (NYD-SP25), mRNA
NM 032231	Homo sapiens hypothetical protein FLJ22875 (FLJ22875), mRNA
NM_018437	Homo sapiens hypothetical protein EDAG-1 (EDAG-1), mRNA
NM_033378	Homo sapiens chorionic gonadotropin, beta polypeptide 2 (CGB2), mRNA
NM_033377	Homo sapiens chorionic gonadotropin, beta polypeptide 1 (CGB1), mRNA
NM_033448	Homo sapiens keratin 6 irs (KRT6IRS), mRNA
NM_033424	Homo sapiens similar to MYOSIN HEAVY CHAIN, CARDIAC MUSCLE
	ALPHA ISOFORM (MYHC-ALPHA) (M. musculus) (LOC92771), mRNA
NM_033445	Homo sapiens similar to H2A histone family, member A (H. sapiens)
	(MGC3165), mRNA
NM_033439	Homo sapiens DVS27-related protein (DVS27), mRNA
NM_033440	Homo sapiens elastase 2A (ELA2A), mRNA
NM_033438	Homo sapiens CD84-H1 precursor (CD84-H1), mRNA
NM_033423	Homo sapiens similar to granzyme B (granzyme 2, cytotoxic T-lymphocyte-
NM 033411	associated serine esterase 1) (H. sapiens) (CTLA1), mRNA
NM_033411	Homo sapiens hypothetical protein MGC13523 (MGC13523), mRNA Homo sapiens similar to HYPOTHETICAL 34.0 KDA PROTEIN ZK795.3 IN
14141_055410	CHROMOSOME IV (MGC19606), mRNA
NM_033413	Homo sapiens hypothetical gene MGC16309 (MGC16309), mRNA
NM 033410	Homo sapiens hypothetical protein MGC13138 (MGC13138), mRNA
NM 033419	Homo sapiens hypothetical gene MGC9753 (MGC9753), mRNA
NM 014083	Homo sapiens PRO0767 protein (PRO0767), mRNA
NM 033043	Homo sapiens chorionic gonadotropin, beta polypeptide 5 (CGB5), mRNA
NM_031451	Homo sapiens hypothetical protein MGC4766 similar to testis specific protein
	TES101RP (MGC4766), mRNA
NM_033183	Homo sapiens chorionic gonadotropin, beta polypeptide 8 (CGB8), mRNA
NM_020443	Homo sapiens hypothetical protein MGC14961 (MGC14961), mRNA
NM_033343	Homo sapiens LIM homeobox protein 4 (LHX4), mRNA
NM_033318	Homo sapiens hypothetical gene supported by AL449243 (LOC91689), mRNA
NM_033328	Homo sapiens capping protein alpha 3 (CAPPA3), mRNA
NM_033315	Homo sapiens ras-like protein VTS58635 (VTS58635), mRNA
NM 033309	Homo sapiens hypothetical protein MGC4655 (MGC4655), mRNA
NM_033296	Homo sapiens T-cell activation protein (PGR1), mRNA
NM_033297	Homo sapiens leucine-rich-repeat protein (RNO2), mRNA
NM_033280	Homo sapiens similar to signal peptidase complex (18kD) (LOC90701), mRNA
NM_033196	Homo sapiens similar to ZINC FINGER PROTEIN 85 (ZINC FINGER
NIM 022272	PROTEIN HPF4) (HTF1) (H. sapiens) (LOC91120), mRNA
NM_033272	Homo sapiens potassium channel subunit HERG-3 (HERG-3), mRNA

NM_033261	Homo sapiens diphosphate dimethylallyl diphosphate isomerase 2 (IDI2), mRNA
NM_033254	Homo sapiens brother of CDO (BOC), mRNA
NM_033204	Homo sapiens hypothetical gene DKFZp570I0164 (DKFZp570I0164), mRNA
NM_033259	Homo sapiens CaM-KII inhibitory protein (CAM-KIIN), mRNA
NM_032597	Homo sapiens testes development-related NYD-SP21 (NYD-SP21), mRNA
NM_033212	Homo sapiens hypothetical gene supported by BC004307; BC008285
	(MGC10992), mRNA
NM_033208	Homo sapiens similar to jerky (mouse) homolog-like (LOC91151), mRNA
NM_033195	Homo sapiens lactate dehydrogenase A -like (LDHL), mRNA
NM_015643	Homo sapiens DKFZP434F122 protein (DKFZP434F122), mRNA
NM_032604	Homo sapiens lung alpha/beta hydrolase 1 (LABH1), mRNA
NM_032133	Homo sapiens hypothetical protein DKFZp434N1415 (DKFZP434N1415),
	mRNA ·
NM_030803	Homo sapiens hypothetical protein FLJ10035 (FLJ10035), mRNA
NM_024062	Homo sapiens hypothetical protein MGC5338 (MGC5338), mRNA
NM_024059	Homo sapiens hypothetical protein MGC5356 (MGC5356), mRNA
NM_016542	Homo sapiens serine/threonine protein kinase MASK (MST4), mRNA
NM_033127	Homo sapiens regucalcin gene promotor region related protein (RGPR), mRNA
NM_033128	Homo sapiens scinderin (SCIN), mRNA
NM_033058	Homo sapiens ring finger protein 29 (RNF29), mRNA
NM_033116	Homo sapiens hypothetical protein MGC16714 (MGC16714), mRNA
NM_033123	Homo sapiens testis-development related NYD-SP27 (NYD-SP27), mRNA
NM_033126	Homo sapiens serine/threonine kinase PSKH2 (PSKH2), mRNA
NM_033124	Homo sapiens NYD-SP28 protein (NYD-SP28), mRNA
NM_033122	Homo sapiens testis development protein NYD-SP26 (NYD-SP26), mRNA
NM_033114	Homo sapiens MADP-1 protein (MADP-1), mRNA
NM_033083	Homo sapiens EAF1 protein (EAF1), mRNA
NM_033087	Homo sapiens hypothetical protein FLJ14511 (FLJ14511), mRNA
NM_024512	Homo sapiens leucine-rich repeat-containing 2 (LRRC2), mRNA
NM_006029	Homo sapiens paraneoplastic antigen MA1 (PNMA1), mRNA
NM_033025	Homo sapiens hypothetical protein FLJ13511 (7h3), mRNA
NM_015169	Homo sapiens homolog of yeast ribosome biogenesis regulatory protein RRS1
	(RRS1), mRNA
NM_015129	Homo sapiens septin 6 (SEP2), mRNA
NM_032838	Homo sapiens hypothetical protein FLJ14779 (FLJ14779), mRNA
NM_032206	Homo sapiens hypothetical protein FLJ21709 (FLJ21709), mRNA
NM_032797	Homo sapiens hypothetical protein FLJ14497 (FLJ14497), mRNA
NM_032472	Homo sapiens peptidylprolyl isomerase (cyclophilin)-like 3 (PPIL3), mRNA
NM_032936	Homo sapiens DC32 (DC32), mRNA
NM_032577	Homo sapiens melanoma-associated chondroitin sulfate proteoglycan-like
17.5.00000	(LOC84664), mRNA
NM_032933	Homo sapiens hypothetical protein MGC11386 (MGC11386), mRNA
NM_032929	Homo sapiens hypothetical protein MGC14793 (MGC14793), mRNA
NM_032928	Homo sapiens hypothetical protein MGC14141 (MGC14141), mRNA
NM_032927	Homo sapiens hypothetical protein MGC13159 (MGC13159), mRNA
NM_032926	Homo sapiens hypothetical protein MGC15737 (MGC15737), mRNA
NM_032921	Homo sapiens hypothetical protein MGC15875 (MGC15875), mRNA
NM_032909	Homo sapiens hypothetical protein MGC14139 (MGC14139), mRNA
NM_032908	Homo sapiens hypothetical protein MGC14407 (MGC14407), mRNA
NM_032906	Homo sapiens hypothetical protein MGC14156 (MGC14156), mRNA
NM_032905	Homo sapiens hypothetical protein MGC14439 (MGC14439), mRNA
NM 032903	Homo sapiens hypothetical protein MGC14425 (MGC14425), mRNA

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NM_032902	Homo sapiens protein phosphatase 1, regulatory (inhibitor) subunit 16A
	(PPP1R16A), mRNA
NM_032901	Homo sapiens hypothetical protein MGC14288 (MGC14288), mRNA
NM_032899	Homo sapiens hypothetical protein MGC14128 (MGC14128), mRNA
NM_032898	Homo sapiens hypothetical protein MGC14126 (MGC14126), mRNA
NM_032897_	Homo sapiens hypothetical protein MGC14436 (MGC14436), mRNA
NM_032896	Homo sapiens hypothetical protein MGC14388 (MGC14388), mRNA
NM_032892	Homo sapiens hypothetical protein MGC14161 (MGC14161), mRNA
NM 032891	Homo sapiens hypothetical protein MGC12928 (MGC12928), mRNA
NM 032890	Homo sapiens hypothetical protein MGC13130 (MGC13130), mRNA
NM 032887	Homo sapiens hypothetical protein MGC16037 (MGC16037), mRNA
NM 032885	Homo sapiens hypothetical protein MGC15906 (MGC15906), mRNA
NM 032882	Homo sapiens hypothetical protein MGC15827 (MGC15827), mRNA
NM 032881	Homo sapiens U7 snRNP-specific Sm-like protein LSM10 (LSM10), mRNA
NM 032880	Homo sapiens hypothetical protein MGC15730 (MGC15730), mRNA
NM 032878	Homo sapiens hypothetical protein MGC15677 (MGC15677), mRNA
NM 032873	Homo sapiens hypothetical protein MGC15437 (MGC15437), mRNA
NM 032867	Homo sapiens hypothetical protein FLJ14966 (FLJ14966), mRNA
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NM 032855	Homo sapiens hematopoietic SH2 protein (HSH2), mRNA
NM 032854	Homo sapiens hypothetical protein FLJ14871 (FLJ14871), mRNA
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NM 032836	Homo sapiens hypothetical protein FLJ14768 (FLJ14768), mRNA
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	(PPP1R15B), mRNA
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NM 032829	Homo sapiens hypothetical protein FLJ14721 (FLJ14721), mRNA
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NM_032727	Homo sapiens internexin neuronal intermediate filament protein, alpha (INA), mRNA
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NM 032709	Homo sapiens hypothetical protein MGC13047 (MGC13047), mRNA
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NM 032627	Homo sapiens hypothetical protein MGC3181 (MGC3181), mRNA
NM 032626	Homo sapiens hypothetical brain protein my038 (MY038), mRNA
NM 032624	Homo sapiens hypothetical brain protein my050 (MY050), mRNA
NM 032623	Homo sapiens ovary-specific acidic protein (OSAP), mRNA
NM_032622	Homo sapiens multi-PDZ-domain-containing protein (LNX), mRNA
NM 032620	Homo sapiens mitochondrial GTP binding protein (GTPBG3), mRNA
NM 018622	Homo sapiens presenilins associated rhomboid-like protein (PARL), mRNA
NM 032498	Homo sapiens homeobox protein from AL590526 (LOC84528), mRNA
NM 032600	Homo sapiens testes development-related NYD-SP17 (NYD-SP17), mRNA
NM_032599	Homo sapiens testes development-related NYD-SP18 (NYD-SP18), mRNA
NM 032594	Homo sapiens insulinoma-associated protein IA-6 (INSM2), mRNA
NM 032585	Homo sapiens testis-specific transcript, Y-linked 6 (TTTY6), mRNA
NM 032575	Homo sapiens Kruppel-like zinc finger protein GLIS2 (GLIS2), mRNA
NM 032573	Homo sapiens testis-specific protein TSP-NY (TSP-NY), mRNA
NM 032572	Homo sapiens ribonuclease 7 (RNASE7), mRNA
NM 032568	Homo sapiens GABA(A) receptors associated protein like 3 (GABARAPL3),
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NM_032567	Homo sapiens testis-specific protein NYD-TSP1 (NYD-TSP1), mRNA
NM_032566	Homo sapiens esophagus cancer-related gene-2 (ECG2), mRNA
NM_032562	Homo sapiens group XIII secreted phospholipase A2 (PLA2G13), mRNA
NM 032547	Homo sapiens short coiled-coil protein (HRIHFB2072), mRNA
NM 032546	Homo sapiens ring finger protein 30 (RNF30), mRNA
NM 032519	Homo sapiens hypothetical protein HT023 (HT023), mRNA
NM_032513	Homo sapiens hypothetical protein MGC11303 similar to Zink transporter 2
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NM 032490	Homo sapiens PNAS-127 protein (PNAS-127), mRNA
NM_032488	Homo sapiens protein related with psoriasis (LOC84518), mRNA
NM_032471	Homo sapiens protein kinase (cAMP-dependent, catalytic) inhibitor beta (PKIB),
NM 032292	mRNA Home conicae hymothetical meetain EU (20202 (EU (20202)
NM 032263	Homo sapiens hypothetical protein FLJ20203 (FLJ20203), mRNA
NM 015178	Homo sapiens hypothetical protein DKFZp434B227 (DKFZp434B227), mRNA Homo sapiens KIAA0717 protein (KIAA0717), mRNA
NM 032410	Homo sapiens hook3 protein (HOOK3), mRNA
NM 032108	Homo sapiens sema domain, transmembrane domain (TM), and cytoplasmic
INIVI_032108	domain, (semaphorin) 6B (SEMA6B), mRNA
NM 015636	Homo sapiens DKFZP586J0119 protein (DKFZP586J0119), mRNA
NM 015701	Homo sapiens hypothetical protein (CL25084), mRNA
NM 015224	Homo sapiens KIAA1105 protein (RAP140), mRNA
NM 032390	Homo sapiens nucleolar protein interacting with the FHA domain of pKi-67
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NM 032388	Homo sapiens nasopharyngeal carcinoma-related protein (NPCR), mRNA
NM 032383	Homo sapiens Hermansky-Pudlak syndrome 3 (HPS3), mRNA
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NM_032319	Homo sapiens hypothetical protein MGC13057 (MGC13057), mRNA Homo sapiens chromosome 2 open reading frame 7 (C2orf7), mRNA
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NM_032315	Homo sapiens hypothetical protein MGC4399 (MGC4399), mRNA
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NM 032207	Homo sapiens hypothetical protein FLJ21742 (FLJ21742), mRNA
NM 032205	Homo sapiens hypothetical protein FLJ21615 (FLJ21615), mRNA
NM 032196	Homo sapiens hypothetical protein KIAA1259 (KIAA1259), mRNA
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NM 032164	Homo sapiens hypothetical protein FLJ12298 (FLJ12298), mRNA
NM 032162	Homo sapiens hypothetical protein FLJ11952 (FLJ11952), mRNA
NM 032155	Homo sapiens hypothetical protein DKFZp547I094 (DKFZp547I094), mRNA
NM 032152	Homo sapiens PRAM-1 protein (PRAM-1), mRNA
NM 032149	Homo sapiens hypothetical protein DKFZp434G072 (DKFZP434G072), mRNA
NM_032147	Homo sapiens hypothetical protein DKFZp434D0127 (DKFZP434D0127), mRNA
NM_032146	Homo sapiens hypothetical protein DKFZp434L1123 similar to mouse Arl6 (DKFZP434L1123), mRNA
NM_032143	Homo sapiens hypothetical protein DKFZp434B1727 (DKFZP434B1727), mRNA
NM 032142	Homo sapiens hypothetical protein FLJ10352 (FLJ10352), mRNA
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NM_032118	Homo sapiens hypothetical protein FLJ12953 similar to Mus musculus D3Mm3e (FLJ12953), mRNA
NM_032117	Homo sapiens GAJ protein (GAJ), mRNA
NM_032116	Homo sapiens hypothetical protein MGC2599 similar to katanin p60 subunit A 1 2599 (MGC2599), mRNA
NM_032112	Homo sapiens mitochondrial ribosomal protein L43 (MRPL43), mRNA
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NM_020726	Homo sapiens neurolysin (metallopeptidase M3 family) (NLN), mRNA
NM_020707	Homo sapiens KIAA1173 protein (KIAA1173), mRNA
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NM 015070 Homo sapiens KIAA0853 protein (KIAA0853), mRNA	NM_015196	Homo sapiens KIAA0922 protein (KIAA0922), mRNA
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NM_001221	NM_032308	Homo sapiens hypothetical protein MGC4189 (MGC4189), mRNA
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mRNA NM 031905 Homo sapiens hypothetical protein MGC3195 (MGC3195), mRNA		
NM 031905 Homo sapiens hypothetical protein MGC3195 (MGC3195), mRNA	NM_031908	
NM_031889 Homo sapiens enamelin (ENAM), mRNA		
· 	NM_031889	Homo sapiens enamelin (ENAM), mRNA

NM_022447	Homo sapiens topoisomerase-related function protein 4-2 (TRF4-2), mRNA
NM_031485	Homo sapiens glutamate rich WD repeat protein GRWD (GRWD), mRNA
NM_031484	Homo sapiens hypothetical protein MGC4415 (MGC4415), mRNA
NM_031479	Homo sapiens hypothetical protein MGC4638 (MGC4638), mRNA
NM_031474	Homo sapiens hypothetical protein DKFZp761G1913 (DKFZP761G1913),
	mRNA
NM_031466	Homo sapiens KIAA1882 protein (MGC4737), mRNA
NM_031465	Homo sapiens hypothetical protein MGC13204 (MGC13204), mRNA
NM_031464	Homo sapiens hypothetical protein MGC11287 similar to ribosomal protein S6 kinase, (MGC11287), mRNA
NM 031459	Homo sapiens sestrin 2 (SES2), mRNA
NM 031455	Homo sapiens hypothetical protein DKFZp761F241 (DKFZP761F241), mRNA
NM 031453	Homo sapiens hypothetical protein MGC11034 (MGC11034), mRNA
NM 031452	Homo sapiens hypothetical protein MGC2560 (MGC2560), mRNA
NM 031449	Homo sapiens KIAA1886 protein (DKFZP761I2123), mRNA
NM 031447	Homo sapiens hypothetical protein MGC13033 (MGC13033), mRNA
NM 031446	Homo sapiens hypothetical protein PNAS-131 (PNAS-131), mRNA
NM 031437	Homo sapiens hypothetical protein MGC10823 (MGC10823), mRNA
NM 031436	Homo sapiens hypothetical protein MGC10612 (MGC10612), mRNA
NM 031435	Homo sapiens hypothetical protein DKFZp564I0422 (DKFZP564I0422), mRNA
NM 031430	Homo sapiens rab interacting lysosomal protein (RILP), mRNA
177	Homo sapiens hypothetical protein MGC10812 (MGC10812), mRNA
NM_031425	Homo sapiens hypothetical protein NUF2R (NUF2R), mRNA
NM_031423	Homo sapiens hypothetical protein INOFZR (NOFZR), indiv. Homo sapiens hypothetical protein DKFZp434H0115 (DKFZP434H0115),
NM_031421	mRNA
NM_031412	Homo sapiens GABA(A) receptor-associated protein like 1 (GABARAPL1), mRNA
NM 004637	Homo sapiens RAB7, member RAS oncogene family (RAB7), mRNA
NM 031283	Homo sapiens HMG-box transcription factor TCF-3 (TCF-3), mRNA
NM 031307	Homo sapiens hypothetical protein FKSG32 (FKSG32), mRNA
NM_031305	Homo sapiens hypothetical protein DKFZp564B1162 (DKFZP564B1162), mRNA
NM_031301	Homo sapiens hypothetical protein DKFZp564D0372 (DKFZP564D0372), mRNA
NM_031298	Homo sapiens hypothetical protein MGC2963 (MGC2963), mRNA
NM 031293	Homo sapiens hypothetical protein DKFZp434G131 (DKFZP434G131), mRNA
NM_031292	Homo sapiens hypothetical protein DKFZp434G1415 (DKFZP434G1415), mRNA
NM 031288	Homo sapiens PAP-1 binding protein (PAPA-1), mRNA
NM 031284	Homo sapiens hypothetical protein DKFZp434B195 (DKFZP434B195), mRNA
NM 030972	Homo sapiens hypothetical protein MGC5384 (MGC5384), mRNA
NM 030901	Homo sapiens olfactory receptor, family 7, subfamily A, member 17 (OR7A17),
14141_030501	mRNA
NM 017990	Homo sapiens hypothetical protein FLJ10079 (FLJ10079), mRNA
NM 031219	Homo sapiens hypothetical protein MGC12904 (MGC12904), mRNA
NM 031218	Homo sapiens hypothetical protein FLJ12488 (FLJ12488), mRNA
NM 031214	Homo sapiens hypothetical protein AF311304 (AF311304), mRNA
NM 031214	Homo sapiens hypothetical protein DC50 (DC50), mRNA
NIN # 021207	Homo caniens hymothetical protein HT036 (HT036) mRNA
NM_031207	Homo sapiens hypothetical protein HT036 (HT036), mRNA Homo sapiens WW domain containing protein 1 (WWP1) mRNA
NM_007013	Homo sapiens WW domain-containing protein 1 (WWP1), mRNA

	subunit 5 (MGC3038), mRNA
NB4 020071	Homo sapiens similar to rat tricarboxylate carrier-like protein (BA108L7.2),
NM_030971	1
AD 6 020065	mRNA
NM_030965	Homo sapiens similar to sialyltransferase 7 ((alpha-N-acetylneuraminyl 2,3-
	betagalactosyl-1,3)-N-acetyl galactosaminide alpha-2,6-sialyltransferase) E
377 6 0000 60	(MGC3184), mRNA
NM_030960	Homo sapiens sperm acrosome associated 1 (SPACA1), mRNA
NM_030958	Homo sapiens organic anion transporter polypeptide-related protein 4
NA 020050	(OATPRP4), mRNA
NM_030952	Homo sapiens hypothetical protein DKFZp434J037 (DKFZP434J037), mRNA
NM_030940	Homo sapiens hypothetical protein MGC4276 similar to CG8198 (MGC4276),
NB (020027	mRNA
NM_030937	Homo sapiens hypothetical protein hCLA-iso (HCLA-ISO), mRNA
NM_030929	Homo sapiens hypothetical protein FKSG28 (FKSG28), mRNA
NM_030921	Homo sapiens hypothetical protein DC42 (DC42), mRNA
NM_030917	Homo sapiens hypothetical protein DKFZp586K0717 (DKFZP586K0717),
NDM 020015	mRNA Home coming hymothetical protein DVE7-5661001 (DVE7D5661001) DNA
NM 030915	Homo sapiens hypothetical protein DKFZp566J091 (DKFZP566J091), mRNA
NM 030914	Homo sapiens hypothetical protein MGC2668 (MGC2668), mRNA
NM_030907	Homo sapiens hypothetical protein MGC10731 (MGC10731), mRNA Homo sapiens hypothetical protein FLJ14129 (FLJ14129), mRNA
NM_030895 NM_030891	· · · · · · · · · · · · · · · · · · ·
	Homo sapiens leucine-rich repeat-containing 3 (LRRC3), mRNA Homo sapiens thioredoxin domain-containing (TXNDC), mRNA
NM_030755	
NM_030819	Homo sapiens hypothetical protein MGC11335 (MGC11335), mRNA
NM 030814	Homo sapiens hypothetical protein GL012 (GL012), mRNA
NM_030810	Homo sapiens hypothetical protein MGC3178 (MGC3178), mRNA
NM_030804	Homo sapiens hypothetical protein DKFZp434E2135 (DKFZP434E2135), mRNA
NM 030794	Homo sapiens hypothetical protein FLJ21007 (FLJ21007), mRNA
NM 030759	Homo sapiens nuclear receptor binding factor-2 (NRBF-2), mRNA
NM 030795	Homo sapiens stathmin-like 4 (STMN4), mRNA
NM 020909	Homo sapiens KIAA1548 protein (KIAA1548), mRNA
NM 018023	Homo sapiens hypothetical protein FLJ10201 (FLJ10201), mRNA
NM 023009	Homo sapiens macrophage myristoylated alanine-rich C kinase substrate
14141_023003	(MACMARCKS), mRNA
NM 025230	Homo sapiens hypthetical protein PRO2389 (PRO2389), mRNA
NM 025222	Homo sapiens hypothetical protein PRO2730 (PRO2730), mRNA
NM_025170	Homo sapiens hypothetical protein FLJ12987 (FLJ12987), mRNA
NM 024681	Homo sapiens hypothetical protein FLJ12242 (FLJ12242), mRNA
NM_024928	Homo sapiens hypothetical protein FLJ22559 (FLJ22559), mRNA
NM_017578	Homo sapiens AKAP-binding sperm protein ropporin (DKFZp434B1222),
	mRNA
NM 030642	Homo sapiens apolipoprotein L, 5 (APOL5), mRNA
NM_024513	Homo sapiens FYVE and coiled-coil domain containing 1 (FYCO1), mRNA
NM_030621	Homo sapiens helicase-moi (KIAA0928), mRNA
NM 030641	Homo sapiens apolipoprotein L, 6 (APOL6), mRNA
NM 025190	Homo sapiens KIAA1641 protein (KIAA1641), mRNA
NM 025040	Homo sapiens hypothetical protein FLJ21941 (FLJ21941), mRNA
NM 030613	Homo sapiens hypothetical protein FLJ21628 (FLJ21628), mRNA
NM_024820	Homo sapiens KIAA1608 protein (KIAA1608), mRNA
NM 018015	Homo sapiens hypothetical protein FLJ10178 (FLJ10178), mRNA
NM 024762	Homo sapiens hypothetical protein FLJ21603 (FLJ21603), mRNA
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NM_024329	Homo sapiens hypothetical protein MGC4342 (MGC4342), mRNA
NM_024087	Homo sapiens DKFZP564L0862 protein (DKFZP564L0862), mRNA
NM_030594	Homo sapiens cytoplasmic polyadenylation element binding protein (CPEB1), mRNA
NM_025084	Homo sapiens hypothetical protein FLJ22795 (FLJ22795), mRNA
NM 025090	Homo sapiens KIAA1453 protein (KIAA1453), mRNA
NM 024939	Homo sapiens hypothetical protein FLJ21918 (FLJ21918), mRNA
NM 024903	Homo sapiens hypothetical protein FLJ14297 (FLJ14297), mRNA
NM_024793	Homo sapiens KIAA0643 protein (KIAA0643), mRNA
NM 024718	Homo sapiens hypothetical protein FLJ10101 (FLJ10101), mRNA
NM 015652	Homo sapiens DKFZP564P1916 protein (DKFZP564P1916), mRNA
NM 025189	Homo sapiens hypothetical protein FLJ13659 (FLJ13659), mRNA
NM 025021	Homo sapiens KIAA0616 protein (KIAA0616), mRNA
NM 025010	Homo sapiens KIAA0795 protein (KIAA0795), mRNA
NM 024894	Homo sapiens hypothetical protein FLJ14075 (FLJ14075), mRNA
NM 024840	Homo sapiens hypothetical protein FLJ13590 (FLJ13590), mRNA
NM_022782	Homo sapiens M-phase phosphoprotein 9 (MPHOSPH9), mRNA
NM 017558	Homo sapiens hypothetical protein DKFZp434L0850 (DKFZp434L0850),
_	mRNA
NM_030580	Homo sapiens hypothetical protein MGC10520 (MGC10520), mRNA
NM 025195	Homo sapiens phosphoprotein regulated by mitogenic pathways (C8FW), mRNA
NM 030581	Homo sapiens hypothetical protein FLJ12270 (FLJ12270), mRNA
NM 030577	Homo sapiens hypothetical protein MGC10993 (MGC10993), mRNA
NM 030576	Homo sapiens hypothetical protein MGC10986 (MGC10986), mRNA
NM 030575	Homo sapiens hypothetical protein MGC10334 (MGC10334), mRNA
NM 030572	Homo sapiens hypothetical protein MGC10946 (MGC10946), mRNA
NM_030571	Homo sapiens hypothetical protein MGC10924 similar to Nedd4 WW-binding protein 5 (MGC10924), mRNA
NM 030569	Homo sapiens hypothetical protein MGC10848 (MGC10848), mRNA
NM 030568	Homo sapiens hypothetical protein MGC10818 (MGC10818), mRNA
NM 030567	Homo sapiens hypothetical protein MGC10772 (MGC10772), mRNA
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NM 025132	Homo sapiens KIAA1638 protein (KIAA1638), mRNA
NM 024668	Homo sapiens hypothetical protein FLJ20288 (FLJ20288), mRNA
NM 024547	Homo sapiens KIAA0467 protein (KIAA0467), mRNA
NM 018418	Homo sapiens hypothetical protein (HSD-3.1), mRNA
NM 025182	Homo sapiens hypothetical protein FLJ11560 (FLJ11560), mRNA
NM_025168	Homo sapiens LAP (leucine-rich repeats and PDZ) and no PDZ protein (LANO), mRNA
NM_025081	Homo sapiens KIAA1305 protein (KIAA1305), mRNA
NM_024750	Homo sapiens leucine-rich repeat-containing 2 (LRRC2), mRNA
NM_025266	Homo sapiens hypothetical protein MGC2780 (MGC2780), mRNA
NM 025265	Homo sapiens hypothetical protein MGC2776 (MGC2776), mRNA
NM 025264	Homo sapiens hypothetical protein MGC2454 (MGC2454), mRNA
NM 025247	Homo sapiens hypothetical protein MGC5601 (MGC5601), mRNA
NM_025246	Homo sapiens hypothetical protein MGC3295 (MGC3295), mRNA
NM 025234	Homo sapiens recombination protein REC14 (REC14), mRNA
NM 025221	Homo sapiens calsenilin-like protein (CALP), mRNA
NM 025207	Homo sapiens hypothetical protein PP591 (PP591), mRNA
NM 025204	Homo sapiens hypothetical protein PP2447 (PP2447), mRNA
NM 025203	Homo sapiens hypothetical protein FLJ21945 (FLJ21945), mRNA
NM 025199	Homo sapiens hypothetical protein FLJ20886 (FLJ20886), mRNA

NM_025197	Homo sapiens hypothetical protein FLJ13660 similar to CDK5 activator-binding
	protein C53 (FLJ13660), mRNA
NM_025187	Homo sapiens hypothetical protein FLJ12076 (FLJ12076), mRNA
NM_025184	Homo sapiens hypothetical protein FLJ22843 (FLJ22843), mRNA
NM_025181	Homo sapiens hypothetical protein FLJ22004 (FLJ22004), mRNA
NM 025163	Homo sapiens hypothetical protein FLJ12768 (FLJ12768), mRNA
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NM 025012	Homo sapiens hypothetical protein FLJ13769 (FLJ13769), mRNA
NM 025009	Homo sapiens hypothetical protein FLJ13621 (FLJ13621), mRNA
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NM_024721	Homo sapiens likely ortholog of mouse zinc finger homeodomain 4 (FLJ20980), mRNA
NM_024713	Homo sapiens hypothetical protein FLJ22557 (FLJ22557), mRNA
NM_024712	Homo sapiens engulfment and cell motility 3 (ced-12 homolog, C. elegans) (ELMO3), mRNA
NM 024711	Homo sapiens hypothetical protein FLJ22690 (FLJ22690), mRNA

	
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-	(FLJ12894), mRNA
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NM 024520	Homo sapiens hypothetical protein FLJ22555 (FLJ22555), mRNA
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NM_024519	Homo sapiens hypothetical protein FLJ13725 (FLJ13725), mRNA
NM 024509	Homo sapiens hypothetical protein MGC2656 (MGC2656), mRNA
NM_024506	Homo sapiens hypothetical protein MGC10771 (MGC10771), mRNA
NM_022893	Homo sapiens B-cell CLL/lymphoma 11A (zinc finger protein) (BCL11A),
	mRNA
NM_015113	Homo sapiens KIAA0399 protein (KIAA0399), mRNA
NM_015545	Homo sapiens KIAA0632 protein (KIAA0632), mRNA
NM_020299	Homo sapiens aldo-keto reductase family 1, member B10 (aldose reductase)
	(AKR1B10), mRNA
NM_003308	Homo sapiens testis specific protein, Y-linked (TSPY), mRNA
NM_024339	Homo sapiens hypothetical protein MGC2655 (MGC2655), mRNA
NM_024334	Homo sapiens hypothetical protein MGC3222 (MGC3222), mRNA
NM_024328	Homo sapiens hypothetical protein MGC2652 (MGC2652), mRNA
NM_024327	Homo sapiens hypothetical protein MGC2508 (MGC2508), mRNA
NM_024323	Homo sapiens hypothetical protein MGC11271 (MGC11271), mRNA
NM_024322	Homo sapiens hypothetical protein MGC11266 (MGC11266), mRNA
NM_024320	Homo sapiens hypothetical protein MGC11242 (MGC11242), mRNA
NM_024319	Homo sapiens hypothetical protein MGC4174 (MGC4174), mRNA
NM_024314	Homo sapiens hypothetical protein MGC4294 (MGC4294), mRNA
NM_024313	Homo sapiens hypothetical protein MGC3731 (MGC3731), mRNA
NM_024310	Homo sapiens hypothetical protein MGC4090 (MGC4090), mRNA
NM_024303	Homo sapiens hypothetical protein MGC4161 (MGC4161), mRNA
NM_024297	Homo sapiens hypothetical protein MGC2941 (MGC2941), mRNA
NM_024293	Homo sapiens hypothetical protein MGC3035 (MGC3035), mRNA
NM_023003	Homo sapiens transmembrane 6 superfamily member 1 (TM6SF1), mRNA
NM_015254	Homo sapiens kinesin family member 13B (KIF13B), mRNA
NM_015127	Homo sapiens Mid-1-related chloride channel 1 (KIAA0761), mRNA
NM_024033	Homo sapiens hypothetical protein MGC5242 (MGC5242), mRNA
NM_024122	Homo sapiens hypothetical protein MGC4825 (MGC4825), mRNA
NM_024121	Homo sapiens hypothetical protein FLJ20979 (FLJ20979), mRNA
NM_024119	Homo sapiens hypothetical protein FLJ11354 (FLJ11354), mRNA
NM_024117	Homo sapiens hypothetical protein MGC2745 (MGC2745), mRNA
NM_024115 NM_024111	Homo sapiens hypothetical protein MGC4309 (MGC4309), mRNA
NM 024111	Homo sapiens hypothetical protein MGC4504 (MGC4504), mRNA
NM 024109	Homo sapiens hypothetical protein MGC2654 (MGC2654), mRNA Homo sapiens hypothetical protein MGC2650 (MGC2650), mRNA
NM 024108	Homo sapiens hypothetical protein MGC2030 (MGC2030), mRNA Homo sapiens hypothetical protein MGC3123 (MGC3123), mRNA
NM_024106	Homo sapiens hypothetical protein MGC2663 (MGC2663), mRNA
NM_024104	Homo sapiens hypothetical protein MGC2003 (MGC2003), mRNA Homo sapiens hypothetical protein MGC2747 (MGC2747), mRNA
NM 024102	Homo sapiens hypothetical protein MGC2747 (MGC2747), mRNA Homo sapiens hypothetical protein MGC2722 (MGC2722), mRNA
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NM 024094	Homo sapiens hypothetical protein MGC5528 (MGC5528), mRNA
NM 024093	Homo sapiens hypothetical protein MGC5529 (MGC5529), mRNA
NM_024090	Homo sapiens hypothetical protein MGC5487 (LCE), mRNA
NM_024086	Homo sapiens hypothetical protein MGC3329 (MGC3329), mRNA
NM_024085	Homo sapiens hypothetical protein FLJ22169 (FLJ22169), mRNA
NM 024080	Homo sapiens hypothetical protein MGC2849 (MGC2849), mRNA
NM 024076	Homo sapiens hypothetical protein MGC2628 (MGC2628), mRNA
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NM 024071	Homo sapiens hypothetical protein MGC2550 (MGC2550), mRNA
NM_024070	Homo sapiens hypothetical protein MGC2463 (MGC2463), mRNA
NM_024069	Homo sapiens hypothetical protein MGC2749 (MGC2749), mRNA
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NM 024068 Homo sapiens hypothetical protein MGC2731 (MGC2731), mRNA NM 024065 Homo sapiens hypothetical protein MGC3062 (MGC3062), mRNA NM 024061 Homo sapiens hypothetical protein MGC5521 (MGC5521), mRNA NM 024058 Homo sapiens hypothetical protein MGC5590 (MGC5590), mRNA NM 024057 Homo sapiens hypothetical protein MGC5585 (MGC5585), mRNA NM 024053 Homo sapiens hypothetical protein MGC861 (MGC861), mRNA NM 024050 Homo sapiens hypothetical protein MGC2594 (MGC2594), mRNA NM 024049 Homo sapiens hypothetical protein MGC5566 (MGC5566), mRNA	
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NM_024048 Homo sapiens hypothetical protein MGC3020 (MGC3020), mRNA	İ
NM_024046 Homo sapiens hypothetical protein MGC8407 (MGC8407), mRNA	$\overline{}$
NM_024045 Homo sapiens nucleolar protein GU2 (GU2), mRNA	
NM_024041 Homo sapiens hypothetical protein MGC3180 (MGC3180), mRNA	
NM_024039 Homo sapiens hypothetical protein MGC2488 (MGC2488), mRNA	
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NM_024028 Homo sapiens hypothetical protein MGC3265 (MGC3265), mRNA	
NM_024027 Homo sapiens hypothetical protein MGC3279 similar to collectins (MGC327	9),
mRNA	
NM_024025 Homo sapiens hypothetical protein MGC1136 (MGC1136), mRNA	
NM 024006 Homo sapiens hypothetical protein IMAGE3455200 (IMAGE3455200), mR1	ĮA
NM_015653 Homo sapiens DKFZP566F0546 protein (DKFZP566F0546), mRNA	
NM_015147 Homo sapiens KIAA0582 protein (KIAA0582), mRNA	
NM_016481 Homo sapiens hypothetical protein (HSPC219), mRNA	
NM_023940 Homo sapiens hypothetical protein MGC2827 (MGC2827), mRNA	
NM_023938 Homo sapiens hypothetical protein MGC2742 (MGC2742), mRNA	
NM_023931 Homo sapiens hypothetical protein MGC2474 (MGC2474), mRNA	
NM_015517 Homo sapiens MBD2 (methyl-CpG-binding protein)-interacting zinc finger	
protein (MIZF), mRNA	
NM_015540 Homo sapiens DKFZP727M111 protein (DKFZP727M111), mRNA	
NM_015043 Homo sapiens KIAA0676 protein (KIAA0676), mRNA	
NM_023934 Homo sapiens hypothetical protein MGC2495 (MGC2495), mRNA	
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synthetase (FLJ12389), mRNA	
NM_023926 Homo sapiens hypothetical protein FLJ12895 (FLJ12895), mRNA	
NM_023924 Homo sapiens hypothetical protein FLJ13441 (FLJ13441), mRNA	
NM_020239 Homo sapiens small protein effector 1 of Cdc42 (SPEC1), mRNA	
NM_012069 Homo sapiens ATPase, (Na+)/K+ transporting, beta 4 polypeptide (ATP1B4)	'>
mRNA	
NM 023112 Homo sapiens hypothetical protein FLJ21916 (FLJ21916), mRNA	
NM 015324 Homo sapiens KIAA0409 protein (KIAA0409), mRNA	
NM 023079 Homo sapiens hypothetical protein FLJ13855 (FLJ13855), mRNA	
NM 023077 Homo sapiens hypothetical protein FLJ12439 (FLJ12439), mRNA	
NM 023075 Homo sapiens hypothetical protein FLJ11585 (FLJ11585), mRNA	
NM 023074 Homo sapiens hypothetical protein FLJ12644 (FLJ12644), mRNA	
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NM_023071 Homo sapiens hypothetical protein FLJ13117 (FLJ13117), mRNA	
NM_012319 Homo sapiens LIV-1 protein, estrogen regulated (LIV-1), mRNA	
NM_023012 Homo sapiens hypothetical protein FLJ11021 similar to splicing factor,	
arginine/serine-rich 4 (FLJ11021), mRNA	
NM_023008 Homo sapiens hypothetical protein FLJ12949 (FLJ12949), mRNA	

NM 022918 Homo sapiens hypothetical protein FLJ22104 (FLJ22104), mRNA NM 022911 Homo sapiens hypothetical protein FLJ12107 (24432), mRNA NM 022902 Homo sapiens hypothetical protein FLJ13110 (FLJ13110), mRNA NM 022903 Homo sapiens hypothetical protein FLJ21303 (FLJ23053), mRNA NM 022904 Homo sapiens hypothetical protein FLJ12057 (FLJ125772), mRNA NM 022905 Homo sapiens hypothetical protein FLJ21302 (FLJ1302), mRNA NM 022808 Homo sapiens bypothetical protein FLJ120302 (FLJ1302), mRNA NM 022841 Homo sapiens hypothetical protein FLJ12094 (FLJ12994), mRNA NM 022842 Homo sapiens hypothetical protein FLJ12994 (FLJ12994), mRNA NM 022843 Homo sapiens hypothetical protein FLJ12215 (FLJ1223017), mRNA NM 022832 Homo sapiens hypothetical protein FLJ2215 (FLJ122515), mRNA NM 022832 Homo sapiens hypothetical protein FLJ21317 (FLJ23017), mRNA NM 022832 Homo sapiens hypothetical protein FLJ21347 (FLJ23147), mRNA NM 022831 Homo sapiens hypothetical protein FLJ21347 (FLJ21347), mRNA NM 022831 Homo sapiens hypothetical protein FLJ21347 (FLJ21347), mRNA NM 022780 Homo sapiens hypothetical protein FLJ23362 (FLJ2362), mRNA NM 022781 Homo sapiens hypothetical protein FLJ23130 (FLJ2343), mRNA NM 022780 Homo sapiens hypothetical protein FLJ21340 (FLJ21343), mRNA NM 022781 Homo sapiens hypothetical protein FLJ213910 (TMRNA NM 022778 Homo sapiens hypothetical protein FLJ213910 (FLJ13910), mRNA NM 022771 Homo sapiens hypothetical protein FLJ121865 (FLJ12863), mRNA NM 022771 Homo sapiens hypothetical protein FLJ121865 (FLJ12391), mRNA NM 022761 Homo sapiens hypothetical protein FLJ2339 (FLJ23399), mRNA NM 022761 Homo sapiens hypothetical protein FLJ2339 (FLJ23339), mRNA NM 022761 Homo sapiens hypothetical protein FLJ2339 (FLJ23339), mRNA NM 022751 Homo sapiens hypothetical protein FLJ2339 (FLJ23339), mRNA NM 022751 Homo sapiens hypothetical protein FLJ2339 (FLJ22339), mRNA NM 022751 Homo sapiens hypothetical protein FLJ23399 (FLJ23339), mRNA NM 022751 Homo sapiens hypothetical protein FLJ2365 (FLJ12865), mRNA NM 022731 Homo sapiens hypothet		
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NM 02281		Homo sapiens hypothetical protein FLJ23053 (FLJ23053), mRNA
NM 02284	NM_022905	Homo sapiens hypothetical protein FLJ12572 (FLJ12572), mRNA
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	L.DMA
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NM_022445	Homo sapiens thiamin pyrophosphokinase 1 (TPK1), mRNA
NM 022495 NM 022494	Homo sapiens hypothetical protein FLJ12799 (FLJ12799), mRNA
NM_022494	Homo sapiens hypothetical protein FLJ21952 (FLJ21952), mRNA Homo sapiens hypothetical protein FLJ12788 (FLJ12788), mRNA
NM 022488	
NM 022480	Homo sapiens PC3-96 protein (PC3-96), mRNA
NM 022474	Homo sapiens hypothetical protein FLJ12587 (FLJ12587), mRNA
NWI_022474	Homo sapiens hypothetical protein FLJ12615 similar to membrane protein, palmitoylated 3 (MAGUK p55 subfamily member 5) (FLJ12615), mRNA
NM_022455	Homo sapiens androgen receptor-associated coregulator 267 (ARA267), mRNA
NM_022452	Homo sapiens hypothetical protein FLJ11618 (FLJ11618), mRNA
NM_022448	Homo sapiens hypothetical protein FLJ21817 similar to Rhoip2 (FLJ21817), mRNA
NM_022373	Homo sapiens hypothetical protein FLJ22313 (FLJ22313), mRNA
NM_022370	Homo sapiens hypothetical protein FLJ21044 similar to Rbig1 (FLJ21044), mRNA
NM_022368	Homo sapiens praja 1 (PJA1), mRNA
NM_022366	Homo sapiens hypothetical protein FLJ23182 (FLJ23182), mRNA
NM_022361	Homo sapiens popeye protein 3 (POP3), mRNA
NM_022360	Homo sapiens human epididymis-specific 3 beta (HE3-BETA), mRNA
NM_022342	Homo sapiens kinesin family member 9 (KIF9), mRNA
NM_022372	Homo sapiens G protein beta subunit-like (GBL), mRNA
NM_022158	Homo sapiens fructosamine-3-kinase (FN3K), mRNA
NM_022137	Homo sapiens secreted modular calcium-binding protein 1 (SMOC1), mRNA
NM_022118	Homo sapiens cutaneous T-cell lymphoma tumor antigen se70-2 (SE70-2), mRNA
NM_022116	Homo sapiens fidgetin-like 1 (FIGNL1), mRNA
NM_022103	Homo sapiens hypothetical zinc finger protein FLJ14011 (FLJ14011), mRNA
NM_022070	Homo sapiens hypothetical protein FLJ22087 (FLJ22087), mRNA
NM_022065	Homo sapiens hypothetical protein FLJ21877 (FLJ21877), mRNA
NM_021970	Homo sapiens mitogen-activated protein kinase kinase 1 interacting protein 1 (MAP2K1IP1), mRNA
NM_019081	Homo sapiens KIAA0430 gene product (KIAA0430), mRNA
NM_021981	Homo sapiens pre-T/NK cell associated protein (1D12A), mRNA
NM_020121	Homo sapiens UDP-glucose ceramide glucosyltransferase-like 2 (UGCGL2), mRNA
NM_006683	Homo sapiens human epididymis-specific 3 alpha (HE3-ALPHA), mRNA
NM_006077	Homo sapiens calcium binding atopy-related autoantigen 1 (CBARA1), mRNA
NM_021934	Homo sapiens hypothetical protein FLJ11773 (FLJ11773), mRNA
NM_021933	Homo sapiens hypothetical protein FLJ12438 (FLJ12438), mRNA
NM_021930	Homo sapiens Rad50-interacting protein 1 (FLJ11785), mRNA
NM_021929	Homo sapiens hypothetical protein FLJ21613 similar to rat corneal wound
	healing related protein (FLJ21613), mRNA
NM_007272	Homo sapiens chymotrypsin C (caldecrin) (CTRC), mRNA
NM_004237	Homo sapiens thyroid hormone receptor interactor 13 (TRIP13), mRNA
NM_003849	Homo sapiens succinate-CoA ligase, GDP-forming, alpha subunit (SUCLG1), mRNA
NM_021648	Homo sapiens KIAA0721 protein (KIAA0721), mRNA
NM_021831	Homo sapiens hypothetical protein FLJ21839 (FLJ21839), mRNA
NM_021827	Homo sapiens hypothetical protein FLJ23514 (FLJ23514), mRNA
NM_021195	Homo sapiens claudin 6 (CLDN6), mRNA
NM_018947	Homo sapiens cytochrome c (HCS), mRNA

NM 021732 Homo sapiens hypothetical protein PP5395 (PP5395), mRNA NM 021730 Homo sapiens hypothetical protein PP1044 (PP1044), mRNA NM 021643 Homo sapiens GS3955 protein (GS3955), mRNA NM 015180 Homo sapiens synaptic nuclei expressed gene 2 (SYNE-2), mRNA NM 021633 Homo sapiens kelch-like protein C3IP1 (C3IP1), mRNA NM 021629 Homo sapiens guanine nucleotide binding protein beta subunit 4 (GNB4), mRNA NM 021626 Homo sapiens sentrin-specific protease (SENP2), mRNA NM 021626 Homo sapiens likely homolog of rat and mouse retinoid-inducible serine carboxypeptidase (RISC), mRNA NM 021622 Homo sapiens pleckstrin homology domain-containing, family A (phosphoinositide binding specific) member 1 (PLEKHA1), mRNA NM 021252 Homo sapiens protein kinase C binding protein 1 (PRKCBP1), mRNA NM 021252 Homo sapiens RAB18, member RAS oncogene family (RAB18), mRNA NM 021253 Homo sapiens interleukin 22 receptor (IL22R), mRNA NM 021254 Homo sapiens epidermal growth factor receptor substrate EPS15R (EPS15R), mRNA NM 021101 Homo sapiens neurogenic differentiation 4 (NEUROD4), mRNA NM 021101 Homo sapiens enhancer of invasion 10 (HEI10), mRNA NM 021178 Homo sapiens serine protease inhibitor, Kazal type, 2 (acrosin-trypsin inhibitor) (SPINK2), mRNA Homo sapiens interferon induced transmembrane protein 2 (1-8D) (IFITM2), mRNA NM 021103 Homo sapiens bone morphogenetic protein 5 (BMP5), mRNA NM 021073 Homo sapiens bone morphogenetic protein 5 (BMP5), mRNA NM 003142 Homo sapiens bone morphogenetic protein 5 (BMP5), mRNA NM 003184 Homo sapiens muscle specific gene (M9), mRNA NM 003234 Homo sapiens muscle specific gene (M9), mRNA NM 013234 Homo sapiens muscle specific gene (M9), mRNA NM 013234 Homo sapiens muscle specific gene (M9), mRNA NM 013234 Homo sapiens muscle specific gene (M9), mRNA NM 013234 Homo sapiens muscle specific gene (M9), mRNA NM 013234 Homo sapiens muscle specific gene (M9), mRNA NM 013234 Homo sapiens muscle specific
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NM012408Homo sapiens protein kinase C binding protein 1 (PRKCBP1), mRNANM021252Homo sapiens RAB18, member RAS oncogene family (RAB18), mRNANM020806Homo sapiens gephyrin (GPHN), mRNANM021258Homo sapiens interleukin 22 receptor (IL22R), mRNANM021235Homo sapiens epidermal growth factor receptor substrate EPS15R (EPS15R), mRNANM021204Homo sapiens E-1 enzyme (MASA), mRNANM021191Homo sapiens neurogenic differentiation 4 (NEUROD4), mRNANM021178Homo sapiens enhancer of invasion 10 (HEI10), mRNANM021127Homo sapiens phorbol-12-myristate-13-acetate-induced protein 1 (PMAIP1), mRNANM021114Homo sapiens serine protease inhibitor, Kazal type, 2 (acrosin-trypsin inhibitor) (SPINK2), mRNANM021103Homo sapiens thymosin, beta 10 (TMSB10), mRNANM001073Homo sapiens bone morphogenetic protein 5 (BMP5), mRNANM003142Homo sapiens Sjogren syndrome antigen B (autoantigen La) (SSB), mRNANM003888Homo sapiens muscle specific gene (M9), mRNANM013234Homo sapiens muscle specific gene (M9), mRNA
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NM_021235 Homo sapiens epidermal growth factor receptor substrate EPS15R (EPS15R), mRNA NM_021204 Homo sapiens E-1 enzyme (MASA), mRNA NM_021191 Homo sapiens neurogenic differentiation 4 (NEUROD4), mRNA NM_021178 Homo sapiens enhancer of invasion 10 (HEI10), mRNA NM_021127 Homo sapiens phorbol-12-myristate-13-acetate-induced protein 1 (PMAIP1), mRNA NM_021114 Homo sapiens serine protease inhibitor, Kazal type, 2 (acrosin-trypsin inhibitor) (SPINK2), mRNA NM_021103 Homo sapiens thymosin, beta 10 (TMSB10), mRNA NM_006435 Homo sapiens interferon induced transmembrane protein 2 (1-8D) (IFITM2), mRNA NM_021073 Homo sapiens bone morphogenetic protein 5 (BMP5), mRNA NM_003142 Homo sapiens Sjogren syndrome antigen B (autoantigen La) (SSB), mRNA NM_003888 Homo sapiens aldehyde dehydrogenase 1 family, member A2 (ALDH1A2), mRNA NM_013234 Homo sapiens muscle specific gene (M9), mRNA
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NM_021020 Homo sapiens leucine zipper, putative tumor suppressor 1 (LZTS1), mRNA
NM_021025 Homo sapiens homeo box 11-like 2 (HOX11L2), mRNA
NM_021003 Homo sapiens protein phosphatase 1A (formerly 2C), magnesium-dependent,
alpha isoform (PPM1A), mRNA
NM_020674 Homo sapiens cytochrome P450 monooxygenase (CYP-M), mRNA
NM_019612 Homo sapiens hypothetical protein R30953_1 (R30953_1), mRNA
NM_020904 Homo sapiens pleckstrin homology domain-containing, family A
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NM_020686 Homo sapiens NPD009 protein (NPD009), mRNA
NM_020684 Homo sapiens NPD007 protein (NPD007), mRNA
NM_020684Homo sapiens NPD007 protein (NPD007), mRNANM_020683Homo sapiens AD026 protein (AD026), mRNA
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NM 020684 Homo sapiens NPD007 protein (NPD007), mRNA NM 020683 Homo sapiens AD026 protein (AD026), mRNA NM 020679 Homo sapiens AD023 protein (AD023), mRNA NM 020677 Homo sapiens HSCARG protein (HSCARG), mRNA
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NM 020684 Homo sapiens NPD007 protein (NPD007), mRNA NM 020683 Homo sapiens AD026 protein (AD026), mRNA NM 020679 Homo sapiens AD023 protein (AD023), mRNA NM 020677 Homo sapiens HSCARG protein (HSCARG), mRNA NM 020675 Homo sapiens AD024 protein (AD024), mRNA NM 020673 Homo sapiens RAB22A, member RAS oncogene family (RAB22A), mRNA NM 020660 Homo sapiens connexin-36 (CX36), mRNA
NM 020684 Homo sapiens NPD007 protein (NPD007), mRNA NM 020683 Homo sapiens AD026 protein (AD026), mRNA NM 020679 Homo sapiens AD023 protein (AD023), mRNA NM 020677 Homo sapiens HSCARG protein (HSCARG), mRNA NM 020675 Homo sapiens AD024 protein (AD024), mRNA NM 020673 Homo sapiens RAB22A, member RAS oncogene family (RAB22A), mRNA

NM 018434	Homo sapiens goliath protein (GP), mRNA
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1111_020437	mRNA
NM 020524	Homo sapiens hematopoietic PBX-interacting protein (HPIP), mRNA
NM 018638	Homo sapiens ethanolamine kinase (EKI1), mRNA
NM 016326	Homo sapiens chemokine-like factor 1 (CKLF1), mRNA
NM 016951	Homo sapiens chemokine-like factor 1 (CKLF1), mRNA
NM_020143	Homo sapiens putatative 28 kDa protein (LOC56902), mRNA
NM 020141	Homo sapiens protein x 013 (AD-020), mRNA
NM 020122	Homo sapiens potassium channel modulatory factor (PCMF), mRNA
NM 018843	Homo sapiens mitochondrial carrier family protein (MCFP), mRNA
NM 018840	Homo sapiens putative Rab5-interacting protein (RIP5), mRNA
NM 016303	Homo sapiens pp21 homolog (LOC51186), mRNA
NM 016300	Homo sapiens cyclic AMP-regulated phosphoprotein, 21 kD (ARPP-21), mRNA
NM 016299	Homo sapiens likely ortholog of mouse heat shock protein, 70 kDa 4
11.12_010255	(LOC51182), mRNA
NM 013259	Homo sapiens neuronal protein (NP25), mRNA
NM_005064	Homo sapiens small inducible cytokine subfamily A (Cys-Cys), member 23
	(SCYA23), mRNA
NM 013260	Homo sapiens transcriptional regulator protein (HCNGP), mRNA
NM_020433	Homo sapiens hypothetical protein LOC57158 (LOC57158), mRNA
NM 020410	Homo sapiens CGI-152 protein (CGI-152), mRNA
NM 020401	Homo sapiens nuclear pore complex protein (NUP107), mRNA
NM 020400	Homo sapiens G protein-coupled receptor 92 (GPR92), mRNA
NM_020397	Homo sapiens CamKI-like protein kinase (LOC57118), mRNA
NM_020388	Homo sapiens CATX-15 protein (CATX-15), mRNA
NM_020386	Homo sapiens HRAS-like suppressor (HRASLS), mRNA
NM_020361	Homo sapiens carboxypeptidase B precursor (CPAH), mRNA
NM_020357	Homo sapiens PEST-containing nuclear protein (pcnp), mRNA
NM_020345	Homo sapiens I-kappa-B-interacting Ras-like protein 1 (KBRAS1), mRNA
NM_020360	Homo sapiens phospholipid scramblase 3 (PLSCR3), mRNA
NM_020348	Homo sapiens cyclin M1 (CNNM1), mRNA
NM_000888	Homo sapiens integrin, beta 6 (ITGB6), mRNA
NM_020181	Homo sapiens myelin proteolipid protein-like protein (PLPL), mRNA
NM_020144	Homo sapiens poly(A) polymerase beta (testis specific) (PAPOLB), mRNA
NM_020202	Homo sapiens Nit protein 2 (NIT2), mRNA
NM_020250	Homo sapiens MOST2 protein (MOST2), mRNA
NM_020237	Homo sapiens MOST-1 protein (MOST-1), mRNA
NM_020234	Homo sapiens x 009 protein (MDS009), mRNA
NM_020128	Homo sapiens nuclear protein double minute 1 (MDM1), mRNA
NM_020169	Homo sapiens latexin protein (LXN), mRNA
NM_020133	Homo sapiens lysophosphatidic acid acyltransferase-delta (LPAAT-delta),
	mRNA
NM_020241	Homo sapiens sema domain, transmembrane domain (TM), and cytoplasmic
) T (000160	domain, (semaphorin) 6B (SEMA6B), mRNA
NM_020163	Homo sapiens semaphorin sem2 (LOC56920), mRNA
NM_020199	Homo sapiens HTGN29 protein (HTGN29), mRNA
NM_020197	Homo sapiens HSKM-B protein (HSKM-B), mRNA
NM_020200	Homo sapiens HHGP protein (HHGP), mRNA
NM_020195	Homo sapiens HCDI protein (HCDI), mRNA
NM_020198	Homo sapiens GK001 protein (GK001), mRNA
NM_020117	Homo sapiens hypothetical protein FLJ10595 (FLJ10595), mRNA

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NM_020119	Homo sapiens hypothetical protein FLB6421 (FLB6421), mRNA
NM_020162	Homo sapiens DEAD/H (Asp-Glu-Ala-Asp/His) box polypeptide 33 (DDX33), mRNA
NM_020215	Homo sapiens hypothetical protein DKFZp761F2014 (DKFZp761F2014), mRNA
NM 020221	Homo sapiens hypothetical protein DKFZp547I224 (DKFZp547I224), mRNA
NM_020217	Homo sapiens hypothetical protein DKFZp547I014 (DKFZp547I014), mRNA
NM 020161	Homo sapiens hypothetical protein DKFZp547H025 (DKFZp547H025), mRNA
NM_020186	Homo sapiens DC11 protein (DC11), mRNA
NM 020205	Homo sapiens cellular zinc finger anti-NF-kappaB Cezanne (CEZANNE),
	mRNA
NM_019887	Homo sapiens second mitochondria-derived activator of caspase (SMAC), mRNA
NM_019892	Homo sapiens phosphatidylinositol (4,5) bisphosphate 5-phosphatase homolog;
	phosphatidylinositol polyphosphate 5-phosphatase type IV (PPI5PIV), mRNA
NM_019885	Homo sapiens cytochrome P450 retinoid metabolizing protein (P450RAI-2), mRNA
NM_019845	Homo sapiens candidate mediator of the p53-dependent G2 arrest (REPRIMO), mRNA
NM 019853	Homo sapiens protein phosphatase 4 regulatory subunit 2 (PPP4R2), mRNA
NM 013301	Homo sapiens protein predicted by clone 23882 (HSU79303), mRNA
NM 013300	Homo sapiens protein predicted by clone 23733 (HSU79274), mRNA
NM 013296	Homo sapiens LGN protein (HSU54999), mRNA
NM 013293	Homo sapiens transformer-2 alpha (htra-2 alpha) (HSU53209), mRNA
NM 013310	Homo sapiens hypothetical protein (AF038169), mRNA
NM 018975	Homo sapiens TRF2-interacting telomeric RAP1 protein (RAP1), mRNA
NM 019082	Homo sapiens putative nucleolar RNA helicase (NOH61), mRNA
NM 019020	Homo sapiens hypothetical protein (FLJ20748), mRNA
NM 019058	Homo sapiens HIF-1 responsive RTP801 (FLJ20500), mRNA
NM 019056	Homo sapiens neuronal protein 17.3 (P17.3), mRNA
NM 019042	Homo sapiens hypothetical protein (FLJ20485), mRNA
NM_019061	Homo sapiens phosphatidylinositol-3 phosphate 3-phosphatase adaptor subunit
ND 6 010006	(3-PAP), mRNA
NM_018986	Homo sapiens hypothetical protein (FLJ20356), mRNA
NM_019034	Homo sapiens ras homolog gene family, member F (in filopodia) (ARHF), mRNA
NM_019062	Homo sapiens hypothetical protein (FLJ20225), mRNA
NM_019038	Homo sapiens hypothetical protein (FLJ11045), mRNA
NM_019044	Homo sapiens hypothetical protein (FLJ10996), mRNA
NM_018180	Homo sapiens DEAD/H (Asp-Glu-Ala-Asp/His) box polypeptide 32 (DDX32), mRNA
NM_019028	Homo sapiens hypothetical protein similar to ankyrin repeat-containing priotein AKR1 (FLJ10852), mRNA
NM_019014	Homo sapiens similar to DNA-directed RNA polymerase I (135 kDa) (Rpo1-2), mRNA
NM 019023	Homo sapiens hypothetical protein (FLJ10640), mRNA
NM 018162	Homo sapiens hypothetical protein FLJ10633 (FLJ10633), mRNA
NM 019067	Homo sapiens hypothetical protein (FLJ10613), mRNA
NM 019057	Homo sapiens hypothetical protein (FLJ10404), mRNA
NM_018846	Homo sapiens SBBI26 protein (SBBI26), mRNA
NM_016483	Homo sapiens hypothetical protein (HSPC226), mRNA
NM 018400	Homo sapiens voltage-gated sodium channel beta-3 subunit (scn3b gene)
T4141 0 T 0 ± 0 0	Andrie Sapiens voltage-gated sodium channel deta-3 subunit (scn3b gene)

	(HSA243396), mRNA
NM_018700	Homo sapiens tripartite motif-containing 36 (TRIM36), mRNA
NM 018547	H mo servines hypothetical materia PRO2064 (PRO2064) PNA
NM 018546	H mo sapiens hypothetical protein PRO2964 (PRO2964), mRNA
NM_018544	Homo sapiens hypothetical protein PRO2958 (PRO2958), mRNA
NM 018634	Homo sapiens hypothetical protein PRO2949 (PRO2949), mRNA
	Homo sapiens hypothetical protein PRO2893 (PRO2893), mRNA
NM_018543	Homo sapiens hypothetical protein PRO2859 (PRO2859), mRNA
NM_018542	Homo sapiens hypothetical protein PRO2834 (PRO2834), mRNA
NM_018538	Homo sapiens erythroblast membrane-associated protein (ERMAP), mRNA
NM_018534	Homo sapiens hypothetical protein PRO2714 (PRO2714), mRNA
NM_018530	Homo sapiens hypothetical protein PRO2521 (PRO2521), mRNA
NM_018627	Homo sapiens hypothetical protein PRO2405 (PRO2405), mRNA
NM_018523	Homo sapiens hypothetical protein PRO2325 (PRO2325), mRNA
NM_018519	Homo sapiens hypothetical protein PRO2266 (PRO2266), mRNA
NM_018517	Homo sapiens hypothetical protein PRO2214 (PRO2214), mRNA
NM 018621	Homo sapiens hypothetical protein PRO2198 (PRO2198), mRNA
NM_018619	Homo sapiens hypothetical protein PRO2133 (PRO2133), mRNA
NM_018618	Homo sapiens hypothetical protein PRO2121 (PRO2121), mRNA
NM 018616	Homo sapiens hypothetical protein PRO2037 (PRO2037), mRNA
NM_018512	Homo sapiens hypothetical protein PRO2015 (PRO2015), mRNA
NM_018610	Homo sapiens hypothetical protein PRO1942 (PRO1942), mRNA
NM_018510	Homo sapiens hypothetical protein PRO1866 (PRO1866), mRNA
NM_018507	Homo sapiens hypothetical protein PRO1843 (PRO1843), mRNA
NM_018606	Homo sapiens hypothetical protein PRO1787 (PRO1787), mRNA
NM_018589	Homo sapiens hypothetical protein PRO1635 (PRO1635), mRNA
NM_018587	Homo sapiens hypothetical protein PRO1617 (PRO1617), mRNA
NM_018503	Homo sapiens hypothetical protein PRO1598 (PRO1598), mRNA
NM_018586	Homo sapiens hypothetical protein PRO1584 (PRO1584), mRNA
NM_018502	Homo sapiens hypothetical protein PRO1580 (PRO1580), mRNA
NM_018603	Homo sapiens hypothetical protein PRO1496 (PRO1496), mRNA
NM_018584	Homo sapiens hypothetical protein PRO1489 (PRO1489), mRNA
NM_018582	Homo sapiens hypothetical protein PRO1483 (PRO1483), mRNA
NM_018602	Homo sapiens DnaJ (Hsp40) homolog, subfamily A, member 4 (DNAJA4),
ND 6 010570	mRNA
NM_018578	Homo sapiens hypothetical protein PRO1257 (PRO1257), mRNA
NM_018576	Homo sapiens hypothetical protein PRO1163 (PRO1163), mRNA
NM 018497	Homo sapiens hypothetical protein PRO1048 (PRO1048), mRNA
NM_018565	Homo sapiens hypothetical protein PRO0899 (PRO0899), mRNA
NM_018562	Homo sapiens hypothetical protein PRO0386 (PRO0386), mRNA
NM_018590	Homo sapiens hypothetical protein PRO0082 (PRO0082), mRNA
NM_018667	Homo sapiens sphingomyelin phosphodiesterase 3, neutral membrane (neutral
NM 017544	sphingomyelinase II) (SMPD3), mRNA
	Homo sapiens transcription factor NRF (NRF), mRNA
NM_018468	Homo sapiens uncharacterized hematopoietic stem/progenitor cells protein
NM 018467	MDS033 (MDS033), mRNA
141AT_01040\	Homo sapiens uncharacterized hematopoietic stem/progenitor cells protein
NM 018464	MDS032 (MDS032), mRNA
T4141_0 T0404	Homo sapiens uncharacterized hematopoietic stem/progenitor cells protein MDS029 (MDS029), mRNA
NM_018688	
NM_018686	Homo sapiens bridging integrator 3 (BIN3), mRNA
NM_018446	Homo sapiens CMP-N-acetylneuraminic acid synthase (CMAS), mRNA
1111 010440	Homo sapiens glycosyltransferase AD-017 (AD-017), mRNA

NM_018416	Homo sapiens FOXJ2 forkhead factor (FHX), mRNA
NM_018407	Homo sapiens putative integral membrane transporter (LC27), mRNA
NM_018472	Homo sapiens uncharacterized hypothalamus protein HT011 (HT011), mRNA
NM_018471	Homo sapiens uncharacterized hypothalamus protein HT010 (HT010), mRNA
NM_018470	Homo sapiens uncharacterized hypothalamus protein HT009 (HT009), mRNA
NM_018469	Homo sapiens uncharacterized hypothalamus protein HT008 (HT008), mRNA
NM_017523	Homo sapiens XIAP associated factor-1 (HSXIAPAF1), mRNA
NM_017514	Homo sapiens SEX gene (HSSEXGENE), mRNA
NM_017512	Homo sapiens rTS beta protein (HSRTSBETA), mRNA
NM_016536	Homo sapiens HSPC059 protein (HSPC059), mRNA
NM_018553	Homo sapiens ELG protein (HSA277841), mRNA
NM_018403	Homo sapiens transcription factor (SMIF gene) (HSA275986), mRNA
NM_018404	Homo sapiens centaurin, alpha 2 (CENTA2), mRNA
NM_018401	Homo sapiens gene for serine/threonine protein kinase (HSA250839), mRNA
NM_017582	Homo sapiens NICE-5 protein (HSA243666), mRNA
NM_018684	Homo sapiens hepatocellular carcinoma-associated antigen 127 (HCA127), mRNA
NM_018477	Homo sapiens uncharacterized hypothalamus protein HARP11 (HARP11), mRNA
NM_018652	Homo sapiens golgin-like protein (GLP), mRNA
NM_017962	Homo sapiens hypothetical protein FLJ20825 (FLJ20825), mRNA
NM_017961	Homo sapiens hypothetical protein FLJ20813 (FLJ20813), mRNA
NM_017960	Homo sapiens hypothetical protein FLJ20808 (FLJ20808), mRNA
NM_017959	Homo sapiens hypothetical protein FLJ20802 (FLJ20802), mRNA
NM_017958	Homo sapiens hypothetical protein FLJ20783 (FLJ20783), mRNA
NM_017957	Homo sapiens epsin 3 (FLJ20778), mRNA
NM_017956	Homo sapiens hypothetical protein FLJ20772 (FLJ20772), mRNA
NM_017950	Homo sapiens hypothetical protein FLJ20753 (FLJ20753), mRNA
NM_017949	Homo sapiens hypothetical protein FLJ20739 (FLJ20739), mRNA
NM_017946	Homo sapiens hypothetical protein FLJ20731 (FLJ20731), mRNA
NM_017953	Homo sapiens hypothetical protein FLJ20729 (FLJ20729), mRNA
NM_017943	Homo sapiens hypothetical protein FLJ20725 (FLJ20725), mRNA
NM_017941	Homo sapiens hypothetical protein FLJ20721 (FLJ20721), mRNA
NM_017938	Homo sapiens hypothetical protein FLJ20716 (FLJ20716), mRNA
NM_017937	Homo sapiens hypothetical protein FLJ20712 (FLJ20712), mRNA
NM_017932	Homo sapiens hypothetical protein FLJ20700 (FLJ20700), mRNA
NM_017929	Homo sapiens hypothetical protein FLJ20695 (FLJ20695), mRNA
NM_017928	Homo sapiens hypothetical protein FLJ20694 (FLJ20694), mRNA
NM_017925	Homo sapiens hypothetical protein FLJ20686 (FLJ20686), mRNA
NM 017920	Homo sapiens hypothetical protein FLJ20654 (FLJ20654), mRNA
NM 017919	Homo sapiens hypothetical protein FLJ20651 (FLJ20651), mRNA
NM 017918	Homo sapiens hypothetical protein FLJ20647 (FLJ20647), mRNA
NM 017917	Homo sapiens hypothetical protein FLJ20644 (FLJ20644), mRNA
NM_017916 NM_017915	Homo sapiens hypothetical protein FLJ20643 (FLJ20643), mRNA Homo sapiens hypothetical protein FLJ20641 (FLJ20641), mRNA
NM_017913	Homo sapiens hypothetical protein FLJ20641 (FLJ20641), mRNA Homo sapiens hypothetical protein FLJ20637 (FLJ20637), mRNA
NM_017912	Homo sapiens hypothetical protein FLJ20637 (FLJ20637), mRNA Homo sapiens hypothetical protein FLJ20627 (FLJ20627), mRNA
NM_017907	Homo sapiens hypothetical protein FLJ20627 (FLJ20627), mRNA
NM 017907	Homo sapiens hypothetical protein FLJ20625 (FLJ20625), mRNA Homo sapiens hypothetical protein FLJ20618 (FLJ20618), mRNA
NM_017901	Homo sapiens two-pore channel 1, homolog (KIAA1169), mRNA
NM_017900	Homo sapiens hypothetical protein FLJ20608 (FLJ20608), mRNA
NM_017899	Homo sapiens hypothetical protein FLJ20607 (TSC), mRNA
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NM 017894 Home sapiens hypothetical protein FLJ20595 (FLJ20595), mRNA NM 017891 Home sapiens sema domain, immunoglobulin domain (Ig), transmembrane domain (TM) and short cytoplasmic domain, (semaphorin) 4G (SEMA4G), mRNA NM 017881 Home sapiens hypothetical protein FLJ20584 (FLJ20584), mRNA NM 017881 Home sapiens hypothetical protein FLJ20568 (FLJ20568), mRNA NM 017876 Home sapiens hypothetical protein FLJ20559 (FLJ20559), mRNA NM 017873 Home sapiens hypothetical protein FLJ20552 (FLJ20552), mRNA NM 017868 Home sapiens hypothetical protein FLJ20533 (FLJ20535), mRNA NM 017866 Home sapiens hypothetical protein FLJ20533 (FLJ20533), mRNA NM 017866 Home sapiens hypothetical protein FLJ20533 (FLJ20533), mRNA NM 017860 Home sapiens hypothetical protein FLJ20519 (FLJ20519), mRNA NM 017860 Home sapiens hypothetical protein FLJ20519 (FLJ20519), mRNA NM 017861 Home sapiens hypothetical protein FLJ20516 (FLJ20516), mRNA NM 017854 Home sapiens hypothetical protein FLJ20514 (FLJ20514), mRNA NM 017854 Home sapiens hypothetical protein FLJ20514 (FLJ20514), mRNA NM 017853 Home sapiens hypothetical protein FLJ20514 (FLJ20511), mRNA NM 017851 Home sapiens hypothetical protein FLJ20511 (FLJ20511), mRNA NM 017843 Home sapiens hypothetical protein FLJ20509 (FLJ20509), mRNA NM 017834 Home sapiens hypothetical protein FLJ20509 (FLJ20509), mRNA NM 017836 Home sapiens hypothetical protein FLJ20509 (FLJ20509), mRNA NM 017836 Home sapiens hypothetical protein FLJ20406 (FLJ20404), mRNA NM 017836 Home sapiens hypothetical protein FLJ20406 (FLJ20440), mRNA NM 017834 Home sapiens hypothetical protein FLJ20446 (FLJ20444), mRNA NM 017834 Home sapiens hypothetical protein FLJ20446 (FLJ20449), mRNA NM 017834 Home sapiens hypothetical protein FLJ20446 (FLJ20449), mRNA NM 017836 Home sapiens hypothetical protein FLJ20446 (FLJ20449), mRNA NM 017817 Home sapiens hypothetical protein FLJ20445 (FLJ20432), mRNA NM 017814 Home sapiens hypothetical p	NIM 017907	Homo sapiens hypothetical protein FLJ20604 (FLJ20604), mRNA
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NM_017803Homo sapiens hypothetical protein FLJ20399 (FLJ20399), mRNANM_017801Homo sapiens hypothetical protein FLJ20396 (FLJ20396), mRNANM_017799Homo sapiens hypothetical protein FLJ20392 (FLJ20392), mRNA		
NM_017801 Homo sapiens hypothetical protein FLJ20396 (FLJ20396), mRNA NM_017799 Homo sapiens hypothetical protein FLJ20392 (FLJ20392), mRNA		
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NM_017766 Homo sapiens hypothetical protein FLJ20321 (FLJ20321), mRNA		
NM_017765 Homo sapiens hypothetical protein FLJ20320 (FLJ20320), mRNA		
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	o sapiens O-linked mannose beta 1,2-N-acetylglucosaminyltransferase 20277), mRNA
	o sapiens hypothetical protein FLJ20275 (FLJ20275), mRNA
	o sapiens hypothetical protein FLJ20258 (FLJ20258), mRNA
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NM 017721 Hom	
	o sapiens hypothetical protein FLJ20241 (FLJ20241), mRNA
	o sapiens hypothetical protein FLJ20211 (FLJ20211), mRNA
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	o sapiens hypothetical protein FLJ20081 (FLJ20081), mRNA
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	o sapiens hypothetical protein FLJ20071 (FLJ20071), mRNA
	o sapiens protein phosphatase 1, regulatory (inhibitor) subunit 9A
(PPP	1R9A), mRNA
NM 017649 Hom	o sapiens cyclin M2 (CNNM2), mRNA

NM_017644	Homo sapiens hypothetical protein FLJ20059 (FLJ20059), mRNA
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NM_017615	Homo sapiens hypothetical protein FLJ20003 (FLJ20003), mRNA
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NM_018393	Homo sapiens hypothetical protein FLJ11336 (FLJ11336), mRNA
NM_018391	Homo sapiens hypothetical protein FLJ11328 (FLJ11328), mRNA
NM_018389	Homo sapiens GDP-fucose transporter 1 (FLJ11320), mRNA
NM_018388	Homo sapiens hypothetical protein FLJ11316 (FLJ11316), mRNA
NM_018386	Homo sapiens hypothetical protein FLJ11305 (FLJ11305), mRNA
NM_018383	Homo sapiens hypothetical protein FLJ11294 (FLJ11294), mRNA
NM_018380	Homo sapiens DEAD/H (Asp-Glu-Ala-Asp/His) box polypeptide 28 (DDX28),
) D C 010070	mRNA
NM 018379	Homo sapiens hypothetical protein FLJ11280 (FLJ11280), mRNA
NM_018376	Homo sapiens hypothetical protein FLJ11275 (FLJ11275), mRNA
NM_018375	Homo sapiens hypothetical protein FLJ11274 (FLJ11274), mRNA
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NM_018365	Homo sapiens hypothetical protein FLJ11222 (FLJ11222), mRNA
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	Homo sapiens hypothetical protein FLJ11200 (FLJ11200), mRNA
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NM 018335	Homo sapiens hypothetical protein FLJ11132 (FLJ11132), mRNA
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NM 018328	Homo sapiens hypothetical protein FLJ11113 (FLJ11113), mRNA
NM_018326	Homo sapiens hypothetical protein FLJ11110 (FLJ11110), mRNA
NM_018324	Homo sapiens hypothetical protein FLJ11106 (FLJ11106), mRNA
NM_018323	Homo sapiens hypothetical protein FLJ11105 (FLJ11105), mRNA
NM_018321	Homo sapiens hypothetical protein FLJ11100 (FLJ11100), mRNA
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NM_018316	Homo sapiens hypothetical protein FLJ11078 (FLJ11078), mRNA
NM_018314	Homo sapiens hypothetical protein FLJ11068 (FLJ11068), mRNA
NM_018309	Homo sapiens hypothetical protein FLJ11046 (FLJ11046), mRNA
NM_018308	Homo sapiens hypothetical protein FLJ11042 (FLJ11042), mRNA
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NM 018306	Homo sapiens hypothetical protein FLJ11036 (FLJ11036), mRNA
NM_018304	Homo sapiens hypothetical protein FLJ11029 (FLJ11029), mRNA
NM 018302	Homo sapiens hypothetical protein FLJ11017 (FLJ11017), mRNA
NM 018299	Homo sapiens hypothetical protein FLJ11011 (FLJ11011), mRNA
NM_018297	Homo sapiens peptide:N-glycanase similar to yeast PNG1 (FLJ11005), mRNA
NM_018296	Homo sapiens hypothetical protein FLJ11004 (FLJ11004), mRNA
NM_018294	Homo sapiens hypothetical protein FLJ10998 (FLJ10998), mRNA
NM 018292	Homo sapiens hypothetical protein FLJ10989 (FLJ10989), mRNA
NM_018289	Homo sapiens hypothetical protein FLJ10979 (FLJ10979), mRNA
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NM_018271	Homo sapiens hypothetical protein FLJ10916 (FLJ10916), mRNA
NM_018264	Homo sapiens hypothetical protein FLJ10900 (FLJ10900), mRNA
NM_018261	Homo sapiens Sec3-like (SEC3), mRNA
NM_018260	Homo sapiens hypothetical protein FLJ10891 (FLJ10891), mRNA
NM_018259	Homo sapiens hypothetical protein FLJ10890 (FLJ10890), mRNA
NM_018250	Homo sapiens hypothetical protein FLJ10871 (FLJ10871), mRNA
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NM_018228	Homo sapiens hypothetical protein FLJ10811 (FLJ10811), mRNA
NM_018227	Homo sapiens hypothetical protein FLJ10808 (FLJ10808), mRNA
NM_018224	Homo sapiens hypothetical protein FLJ10803 (FLJ10803), mRNA
NM_018222	Homo sapiens parvin, alpha (PARVA), mRNA
NM_018221	Homo sapiens chromosome 2 open reading frame 6 (C2orf6), mRNA
NM_018216	Homo sapiens hypothetical protein FLJ10782 (FLJ10782), mRNA
NM_018215	Homo sapiens hypothetical protein FLJ10781 (FLJ10781), mRNA
NM_018214	Homo sapiens LAP (leucine-rich repeats and PDZ) and no PDZ protein (LANO), mRNA
NM_018210	Homo sapiens hypothetical protein FLJ10769 (FLJ10769), mRNA
NM_018208	Homo sapiens hypothetical protein FLJ10761 (FLJ10761), mRNA
NM_018203	Homo sapiens hypothetical protein FLJ10748 (FLJ10748), mRNA
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NM_018199	Homo sapiens hypothetical protein FLJ10738 (FLJ10738), mRNA
NM_018198	Homo sapiens hypothetical protein FLJ10737 (FLJ10737), mRNA
NM_018196	Homo sapiens epsilon-trimethyllysine hydroxylase (FLJ10727), mRNA
NM_018195	Homo sapiens hypothetical protein FLJ10726 (FLJ10726), mRNA
NM_018190	Homo sapiens hypothetical protein FLJ10715 (FLJ10715), mRNA
NM_018189	Homo sapiens hypothetical protein FLJ10713 (FLJ10713), mRNA

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NM_018183	Homo sapiens hypothetical protein FLJ10701 (FLJ10701), mRNA
NM_018182	Homo sapiens hypothetical protein FLJ10700 (FLJ10700), mRNA
NM_018181	Homo sapiens hypothetical protein FLJ10697 (FLJ10697), mRNA
NM_018176	Homo sapiens hypothetical protein FLJ10675 (FLJ10675), mRNA
NM_018174	Homo sapiens chromosome 19 open reading frame 5 (C19orf5), mRNA
NM_018173	Homo sapiens hypothetical protein FLJ10665 (FLJ10665), mRNA
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NM_018115	Homo sapiens hypothetical protein FLJ10498 (FLJ10498), mRNA
NM_018113	Homo sapiens lipocalin-interacting membrane receptor (LIMR), mRNA
NM_018111	Homo sapiens hypothetical protein FLJ10490 (FLJ10490), mRNA
NM_018110	Homo sapiens hypothetical protein FLJ10488 (FLJ10488), mRNA
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NM_018104	Homo sapiens hypothetical protein FLJ10474 (FLJ10474), mRNA
NM_018096	Homo sapiens hypothetical protein similar to beta-transducin family (FLJ10458),
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NM_018095	Homo sapiens hypothetical protein FLJ10450 (FLJ10450), mRNA
NM_018089	Homo sapiens hypothetical protein FLJ10415 (FLJ10415), mRNA
NM_018088	Homo sapiens hypothetical protein FLJ10408 (FLJ10408), mRNA
NM_018084	Homo sapiens hypothetical protein FLJ10392 (FLJ10392), mRNA
NM_018083	Homo sapiens zinc finger protein 358 (ZNF358), mRNA
NM_018082	Homo sapiens hypothetical protein FLJ10388 (FLJ10388), mRNA
NM_018081	Homo sapiens hypothetical protein FLJ10385 (FLJ10385), mRNA
NM_018080	Homo sapiens hypothetical protein FLJ10381 (FLJ10381), mRNA
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NM_018077	Homo sapiens hypothetical protein FLJ10377 (FLJ10377), mRNA
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NM_018068	Homo sapiens likely ortholog of mouse piwi like homolog 1 (Drosophila)-like
\	(FLJ10351), mRNA
NM_018067	Homo sapiens hypothetical protein FLJ10350 (FLJ10350), mRNA
NM_018066	Homo sapiens hypothetical protein FLJ10349 (FLJ10349), mRNA
NM_018065	Homo sapiens hypothetical protein FLJ10346 (FLJ10346), mRNA
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NM_018014	Homo sapiens B-cell CLL/lymphoma 11A (zinc finger protein) (BCL11A), mRNA
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NM_018012	Homo sapiens hypothetical protein FLJ10157 (FLJ10157), mRNA
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NM_018423	Homo sapiens hypothetical protein DKFZp761P1010 (DKFZp761P1010), mRNA
NM_017597	Homo sapiens hypothetical protein DKFZp761K1824 (DKFZp761K1824), mRNA
NM_018422	Homo sapiens hypothetical protein DKFZp761K1423 (DKFZp761K1423),
ND 6 010401	mRNA
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NM_017594	Homo sapiens hypothetical protein DKFZp761C07121 (DKFZp761C07121), mRNA
NM_017535	Homo sapiens hypothetical protein DKFZp566H0824 (DKFZp566H0824), mRNA
NM 018705	Homo sapiens hypothetical protein DKFZp547G183 (DKFZp547G183), mRNA
NM 017604	Homo sapiens KIAA1023 protein (KIAA1023), mRNA
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NM_017559	Homo sapiens hypothetical protein DKFZp434H2215 (DKFZp434H2215), mRNA
NM_017598	Homo sapiens hypothetical protein DKFZp434C0923 (DKFZp434C0923), mRNA
NM_017577	Homo sapiens hypothetical protein DKFZp434C0328 (DKFZp434C0328), mRNA
NM 014612	Homo sapiens C9orf10 protein (C9orf10), mRNA
NM 018460	Homo sapiens uncharacterized bone marrow protein BM046 (BM046), mRNA
NM 018459	Homo sapiens uncharacterized bone marrow protein BM045 (BM045), mRNA
NM_018451	Homo sapiens centrosomal P4.1-associated protein (CPAP), mRNA
NM 018450	Homo sapiens uncharacterized bone marrow protein BM029 (BM029), mRNA
NM 018674	Homo sapiens putative acid-sensing ion channel (ASIC4), mRNA
NM_017435	Homo sapiens solute carrier family 21 (organic anion transporter), member 14
11112_017 155	(SLC21A14), mRNA
NM 016848	Homo sapiens neuronal Shc (SHC3), mRNA
NM 017432	Homo sapiens prostate tumor over expressed gene 1 (PTOV1), mRNA
NM 016953	Homo sapiens phosphodiesterase 11A (PDE11A), mRNA
NM 013242	Homo sapiens similar to mouse Glt3 or D. malanogaster transcription factor IIB
	(AF093680), mRNA
NM 016267	Homo sapiens TONDU (TONDU), mRNA
NM 015859	Homo sapiens general transcription factor IIA, 1 (37kD and 19kD subunits)
	(GTF2A1), mRNA
NM 016271	Homo sapiens STRIN protein (STRIN), mRNA
NM 016584	Homo sapiens interleukin 23, alpha subunit p19 (IL23A), mRNA
NM 016329	Homo sapiens RU1 (RU1), mRNA
NM 016337	Homo sapiens RNB6 (RNB6), mRNA
NM_016146	Homo sapiens PTD009 protein (PTD009), mRNA
NM_016145	Homo sapiens PTD008 protein (PTD008), mRNA
NM_016144	Homo sapiens PTD002 protein (PTD002), mRNA
NM_016147	Homo sapiens protein phosphatase methylesterase-1 (PME-1), mRNA
NM_016445	Homo sapiens pleckstrin 2 (mouse) homolog (PLEK2), mRNA
NM_016170	Homo sapiens NCX protein (NCX), mRNA
NM_016132	Homo sapiens myelin gene expression factor 2 (MEF-2), mRNA
NM_016586	Homo sapiens MBIP protein (MBIP), mRNA
NM_016547	Homo sapiens calcium binding protein Cab45 precursor (Cab45), mRNA
NM_016530	Homo sapiens RAB-8b protein (LOC51762), mRNA
NM_016442	Homo sapiens type 1 tumor necrosis factor receptor shedding aminopeptidase regulator (ARTS-1), mRNA
NM_016438	Homo sapiens CLST 11240 protein (CLST11240), mRNA
NM_016340	Homo sapiens rap guanine nucleotide exchange factor (RA-GEF-2), mRNA
NM_016306	Homo sapiens DnaJ (Hsp40) homolog, subfamily B, member 11 (DNAJB11), mRNA
NM_016292	Homo sapiens heat shock protein 75 (TRAP1), mRNA
NM_016248	Homo sapiens A kinase (PRKA) anchor protein 11 (AKAP11), mRNA
NM_016207	Homo sapiens cleavage and polyadenylation specific factor 3, 73kD subunit (CPSF3), mRNA
NM_016163	Homo sapiens vesicle transport-related protein (RA410), mRNA
NM_016106	Homo sapiens vesicle transport-related protein (RA410), mRNA
NM_016081	Homo sapiens palladin (KIAA0992), mRNA
NM_015934	Homo sapiens nucleolar protein NOP5/NOP58 (NOP5/NOP58), mRNA
NM_015925	Homo sapiens liver-specific bHLH-Zip transcription factor (LISCH7), mRNA
NM_015878	Homo sapiens ornithine decarboxylase antizyme inhibitor (OAZIN), mRNA
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NM_016284	Homo sapiens KIAA1007 protein (KIAA1007), mRNA
NM_016645	Homo sapiens mesenchymal stem cell protein DSC92 (NEUGRIN), mRNA
NM_016631	Homo sapiens chromosome 21 open reading frame 66 (C21 or 66), mRNA
NM_016576	Homo sapiens GMPR2 for guanosine monophosphate reductase isolog (LOC51292), mRNA
NM_016501	Homo sapiens hypothetical protein FLJ10597 (FLJ10597), mRNA
NM 016500	Homo sapiens hypothetical protein (LOC51260), mRNA
NM_016487	Homo sapiens HSPC230 gene (HSPC230), mRNA
NM 016480	Homo sapiens PABP-interacting protein 2 (PAIP2), mRNA
NM 016433	Homo sapiens glycolipid transfer protein (GLTP), mRNA
NM_016369	Homo sapiens claudin 18 (CLDN18), mRNA
NM 016359	Homo sapiens nucleolar protein ANKT (ANKT), mRNA
NM_016246	Homo sapiens retinal short-chain dehydrogenase/reductase retSDR3
1441_010240	(LOC51171), mRNA
NM_016186	Homo sapiens serine (or cysteine) proteinase inhibitor, clade A (alpha-1
	antiproteinase, antitrypsin), member 10 (SERPINA10), mRNA
NM_016180	Homo sapiens AIM-1 protein (MATP), mRNA
NM_016176	Homo sapiens calcium binding protein Cab45 precursor (Cab45), mRNA
NM 016174	Homo sapiens cerebral cell adhesion molecule (LOC51148), mRNA
NM 016131	Homo sapiens RAB10, member RAS oncogene family (RAB10), mRNA
NM 016031	Homo sapiens elongation of very long chain fatty acids (FEN1/Elo2, SUR4/Elo3,
-	yeast)-like 1 (ELOVL1), mRNA
NM_015955	Homo sapiens C21orf19-like protein (LOC51072), mRNA
NM 015931	Homo sapiens fls485 (LOC51066), mRNA
NM_015879	Homo sapiens sialyltransferase 8C (alpha2,3Galbeta1,4GlcNAcalpha 2,8-
	sialyltransferase) (SIAT8C), mRNA
NM_016368	Homo sapiens myo-inositol 1-phosphate synthase A1 (ISYNA1), mRNA
NM_016488	Homo sapiens hypothetical protein (HSPC232), mRNA
NM_016478	Homo sapiens hypothetical protein (HSPC216), mRNA
NM_016463	Homo sapiens hypothetical protein (HSPC195), mRNA
NM_016410	Homo sapiens hypothetical protein HSPC177 (HSPC177), mRNA
NM_016406	Homo sapiens hypothetical protein (HSPC155), mRNA
NM_016401	Homo sapiens hypothetical protein (HSPC138), mRNA
NM_016400	Homo sapiens Huntingtin interacting protein K (HYPK), mRNA
NM_016396	Homo sapiens hypothetical protein (HSPC129), mRNA
NM_016391	Homo sapiens hypothetical protein (HSPC111), mRNA
NM_015933	Homo sapiens hypothetical protein (HSPC016), mRNA
NM_015932	Homo sapiens hypothetical protein (HSPC014), mRNA
NM_016172	Homo sapiens putative glialblastoma cell differentiation-related (GDBR1), mRNA
NM_016194	Homo sapiens guanine nucleotide binding protein (G protein), beta 5 (GNB5),
NM 016106	mRNA
NM 016196	Homo sapiens KIAA0682 gene product (KIAA0682), mRNA
NM 016553	Homo sapiens nucleoporin 62kD (NUP62), mRNA
NM_016195	Homo sapiens M-phase phosphoprotein 1 (MPHOSPH1), mRNA
NM_016550	Homo sapiens HeLa cyclin-dependent kinase 2 interacting protein (CINP), mRNA
NM_016623	Homo sapiens hypothetical protein (BM-009), mRNA
NM_016237	Homo sapiens anaphase promoting complex subunit 5 (ANAPC5), mRNA
NM_016108	Homo sapiens androgen induced protein (AIG-1), mRNA
NM_014886	Homo sapiens hypothetical protein (YR-29), mRNA
NM_014035	Homo sapiens SBBI31 protein (SBBI31), mRNA
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T-2	
NM_014868	Homo sapiens ring finger protein 10 (RNF10), mRNA
NM_014092	Homo sapiens PRO1575 protein (PRO1575), mRNA
NM_014138	Homo sapiens PRO0659 protein (PRO0659), mRNA
NM_014135	Homo sapiens PRO0641 protein (PRO0641), mRNA
NM_014134	Homo sapiens PRO0628 protein (PRO0628), mRNA
NM_014133	Homo sapiens PRO0618 protein (PRO0618), mRNA
NM_014076	Homo sapiens PRO0611 protein (PRO0611), mRNA
NM_014074	Homo sapiens PRO0529 protein (PRO0529), mRNA
NM_014129	Homo sapiens PRO0478 protein (PRO0478), mRNA
NM_014126	Homo sapiens PRO0365 protein (PRO0365), mRNA
NM_014124	Homo sapiens PRO0255 protein (PRO0255), mRNA
NM_014121	Homo sapiens PRO0233 protein (PRO0233), mRNA
NM_014120	Homo sapiens PRO0214 protein (PRO0214), mRNA
NM_014118	Homo sapiens PRO0159 protein (PRO0159), mRNA
NM_014117	Homo sapiens PRO0149 protein (PRO0149), mRNA
NM_014116	Homo sapiens PRO0132 protein (PRO0132), mRNA
NM_015364	Homo sapiens MD-2 protein (MD-2), mRNA
NM_014020	Homo sapiens LR8 protein (LR8), mRNA
NM_014931	Homo sapiens KIAA1115 protein (KIAA1115), mRNA
NM_014901	Homo sapiens KIAA1100 protein (KIAA1100), mRNA
NM_014908	Homo sapiens KIAA1094 protein (KIAA1094), mRNA
NM_014906	Homo sapiens KIAA1072 protein (KIAA1072), mRNA
NM_014932	Homo sapiens neuroligin 1 (NLGN1), mRNA
NM_014894	Homo sapiens KIAA1056 protein (KIAA1056), mRNA
NM_014956	Homo sapiens KIAA1052 protein (KIAA1052), mRNA
NM_014928	Homo sapiens KIAA1046 protein (KIAA1046), mRNA
NM_014909	Homo sapiens KIAA1036 protein (KIAA1036), mRNA
NM_014939	Homo sapiens KIAA1012 protein (KIAA1012), mRNA
NM_014895	Homo sapiens KIAA1009 protein (KIAA1009), mRNA
NM_014960	Homo sapiens KIAA1001 protein (KIAA1001), mRNA
NM_014950	Homo sapiens KIAA0997 protein (KIAA0997), mRNA
NM_014934	Homo sapiens zinc-finger protein DZIP1 (DZIP1), mRNA
NM_014023	Homo sapiens KIAA0982 protein (KIAA0982), mRNA
NM 014900	Homo sapiens KIAA0977 protein (KIAA0977), mRNA
NM 014929	Homo sapiens KIAA0971 protein (KIAA0971), mRNA
NM 014935	Homo sapiens phosphoinositol 3-phosphate-binding protein-2 (PEPP3), mRNA
NM 014937	Homo sapiens Sac domain-containing inositol phosphatase 2 (SAC2), mRNA
NM 014902	Homo sapiens KIAA0964 protein (KIAA0964), mRNA
NM_014898	Homo sapiens KIAA0961 protein (KIAA0961), mRNA
NM_014942	Homo sapiens ankyrin repeat domain 6 (ANKRD6), mRNA
NM_014959	Homo sapiens tumor up-regulated CARD-containing antagonist of caspase nine
_	(TUCAN), mRNA
NM 014952	Homo sapiens KIAA0945 protein (KIAA0945), mRNA
NM_014904	Homo sapiens KIAA0941 protein (Rab11-FIP2), mRNA
NM_014903	Homo sapiens KIAA0938 protein (KIAA0938), mRNA
NM 014897	Homo sapiens KIAA0924 protein (KIAA0924), mRNA
NM 014883	Homo sapiens KIAA0914 gene product (KIAA0914), mRNA
NM 014949	Homo sapiens KIAA0907 protein (KIAA0907), mRNA
NM 014896	Homo sapiens KIAA0894 protein (KIAA0894), mRNA
NM 014969	Homo sapiens KIAA0893 protein (KIAA0893), mRNA
NM 014966	Homo sapiens DEAD/H (Asp-Glu-Ala-Asp/His) box polypeptide 30 (DDX30),
-	mRNA

NM_015377	Homo sapiens KIAA0889 protein (KIAA0889), mRNA
NM_014936	Homo sapiens ectonucleotide pyrophosphatase/phosphodiesterase 4 (putative
	function) (ENPP4), mRNA
NM_014940	Homo sapiens KIAA0872 protein (KIAA0872), mRNA
NM_014943	Homo sapiens KIAA0854 protein (KIAA0854), mRNA
NM_014926	Homo sapiens KIAA0848 protein (KIAA0848), mRNA
NM_014945	Homo sapiens KIAA0843 protein (KIAA0843), mRNA
NM_014924	Homo sapiens KIAA0831 protein (KIAA0831), mRNA
NM_014703	Homo sapiens KIAA0800 gene product (KIAA0800), mRNA
NM_014650	Homo sapiens KIAA0798 gene product (KIAA0798), mRNA
NM_014660	Homo sapiens KIAA0783 gene product (KIAA0783), mRNA
NM_014726	Homo sapiens KIAA0775 gene product (KIAA0775), mRNA
NM_014690	Homo sapiens KIAA0773 gene product (KIAA0773), mRNA
NM_014805	Homo sapiens KIAA0766 gene product (KIAA0766), mRNA
NM_014869	Homo sapiens KIAA0763 gene product (KIAA0763), mRNA
NM_014804	Homo sapiens KIAA0753 gene product (KIAA0753), mRNA
NM_014632	Homo sapiens KIAA0750 gene product (KIAA0750), mRNA
NM_014796	Homo sapiens KIAA0748 gene product (KIAA0748), mRNA
NM_014719	Homo sapiens KIAA0738 gene product (KIAA0738), mRNA
NM_014828	Homo sapiens KIAA0737 gene product (KIAA0737), mRNA
NM_014849	Homo sapiens likely ortholog of mouse synaptic vesicle glycoprotein 2a (SV2),
	mRNA .
NM_014848	Homo sapiens synaptic vesicle protein 2B homolog (SV2B), mRNA
NM_014718	Homo sapiens KIAA0726 gene product (KIAA0726), mRNA
NM_014652	Homo sapiens importin 13 (IMP13), mRNA
NM_014867	Homo sapiens KIAA0711 gene product (KIAA0711), mRNA
NM_014852	Homo sapiens KIAA0682 gene product (KIAA0682), mRNA
NM_014663	Homo sapiens KIAA0677 gene product (KIAA0677), mRNA
NM_014648	Homo sapiens KIAA0675 gene product (KIAA0675), mRNA
NM_014779	Homo sapiens KIAA0669 gene product (KIAA0669), mRNA
NM_014811	Homo sapiens KIAA0649 gene product (KIAA0649), mRNA
NM_014817	Homo sapiens KIAA0644 gene product (KIAA0644), mRNA
NM_015046	Homo sapiens KIAA0625 protein (KIAA0625), mRNA
NM_014694	Homo sapiens KIAA0605 gene product (KIAA0605), mRNA
NM_014832	Homo sapiens KIAA0603 gene product (KIAA0603), mRNA
NM_014749	Homo sapiens KIAA0586 gene product (KIAA0586), mRNA
NM_014668	Homo sapiens KIAA0575 gene product (KIAA0575), mRNA
NM_014709	Homo sapiens KIAA0570 gene product (KIAA0570), mRNA
NM_014704	Homo sapiens KIAA0562 gene product (KIAA0562), mRNA
NM_014790	Homo sapiens KIAA0555 gene product (KIAA0555), mRNA
NM_014731	Homo sapiens KIAA0552 gene product (KIAA0552), mRNA
NM_014793	Homo sapiens KIAA0547 gene product (KIAA0547), mRNA
NM_014825	Homo sapiens chromosome 21 open reading frame 108 (C21orf108), mRNA
NM_014840	Homo sapiens KIAA0537 gene product (KIAA0537), mRNA
NM_014682	Homo sapiens KIAA0535 gene product (KIAA0535), mRNA
NM_014851	Homo sapiens KIAA0469 gene product (KIAA0469), mRNA
NM 014638	Homo sapiens KIAA0450 gene product (KIAA0450), mRNA
NM_015556	Homo sapiens KIAA0440 protein (KIAA0440), mRNA
NM_014801	Homo sapiens KIAA0435 gene product (KIAA0435), mRNA
NM_014772	Homo sapiens KIAA0427 gene product (KIAA0427), mRNA
NM_014631	Homo sapiens KIAA0418 gene product (KIAA0418), mRNA
NM_014702	
NM_014702	Homo sapiens KIAA0408 gene product (KIAA0408), mRNA

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NM_014672	Homo sapiens KIAA0391 gene product (KIAA0391), mRNA
NM_014717	Homo sapiens KIAA0390 gene product (KIAA0390), mRNA
NM_014686	Homo sapiens KIAA0355 gene product (KIAA0355), mRNA
NM_014872	Homo sapiens KIAA0354 gene product (KIAA0354), mRNA
NM_014830	Homo sapiens KIAA0352 gene product (KIAA0352), mRNA
NM_014636	Homo sapiens Ral guanine nucleotide exchange factor RalGPS1A
	(RALGPS1A), mRNA
NM_014635	Homo sapiens KIAA0336 gene product (KIAA0336), mRNA
NM_014803	Homo sapiens KIAA0335 gene product (KIAA0335), mRNA
NM_014844	Homo sapiens KIAA0329 gene product (KIAA0329), mRNA
NM_014821	Homo sapiens KIAA0317 gene product (KIAA0317), mRNA
NM_014699	Homo sapiens KIAA0296 gene product (KIAA0296), mRNA
NM_014742	Homo sapiens KIAA0255 gene product (KIAA0255), mRNA
NM_014734	Homo sapiens KIAA0247 gene product (KIAA0247), mRNA
NM_014760	Homo sapiens KIAA0218 gene product (KIAA0218), mRNA
NM_014735	Homo sapiens KIAA0215 gene product (KIAA0215), mRNA
NM_014630	Homo sapiens KIAA0211 gene product (KIAA0211), mRNA
NM_014744	Homo sapiens KIAA0210 gene product (KIAA0210), mRNA
NM_014725	Homo sapiens KIAA0189 gene product (KIAA0189), mRNA
NM_014753	Homo sapiens KIAA0187 gene product (KIAA0187), mRNA
NM_014791	Homo sapiens likely ortholog of maternal embryonic leucine zipper kinase
	(KIAA0175), mRNA
NM_014746	Homo sapiens KIAA0161 gene product (KIAA0161), mRNA
NM_014633	Homo sapiens KIAA0155 gene product (KIAA0155), mRNA
NM_014002	Homo sapiens IKK-related kinase epsilon; inducible IkappaB kinase (IKKE),
	mRNA
NM_014847	Homo sapiens KIAA0144 gene product (KIAA0144), mRNA
NM_014773	Homo sapiens KIAA0141 gene product (KIAA0141), mRNA
NM_014649	Homo sapiens KIAA0138 gene product (KIAA0138), mRNA
NM_014792	Homo sapiens KIAA0125 gene product (KIAA0125), mRNA
NM_014999	Homo sapiens KIAA0118 protein (KIAA0118), mRNA
NM_014740	Homo sapiens KIAA0111 gene product (KIAA0111), mRNA
NM_014673	Homo sapiens KIAA0103 gene product (KIAA0103), mRNA
NM_014736	Homo sapiens KIAA0101 gene product (KIAA0101), mRNA
NM_014669	Homo sapiens KIAA0095 gene product (KIAA0095), mRNA
NM_014679	Homo sapiens KIAA0092 gene product (KIAA0092), mRNA
NM_014769	Homo sapiens KIAA0087 gene product (KIAA0087), mRNA
NM_014877	Homo sapiens helicase KIAA0054 (KIAA0054), mRNA
NM_014716	Homo sapiens centaurin, beta 1 (CENTB1), mRNA
NM_015361	Homo sapiens R3H domain (binds single-stranded nucleic acids) containing
	(R3HDM), mRNA
NM_014880	Homo sapiens KIAA0022 gene product (KIAA0022), mRNA
NM_014878	Homo sapiens KIAA0020 gene product (KIAA0020), mRNA
NM_014665	Homo sapiens KIAA0014 gene product (KIAA0014), mRNA
NM_014671	Homo sapiens ubiquitin-protein isopeptide ligase (E3) (KIAA0010), mRNA
NM_014637	Homo sapiens KIAA0009 gene product (KIAA0009), mRNA
NM_015384	Homo sapiens IDN3 protein (IDN3), mRNA
NM_014188	Homo sapiens HSPC182 protein (HSPC182), mRNA
NM_014187	Homo sapiens HSPC171 protein (HSPC171), mRNA
NM_014182	Homo sapiens HSPC160 protein (HSPC160), mRNA
NM_014178	Homo sapiens HSPC156 protein (HSPC156), mRNA
NM_014177	Homo sapiens HSPC154 protein (HSPC154), mRNA
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NM 014176	Homo sapiens HSPC150 protein similar to ubiquitin-conjugating enzyme
1411/0	
ND4 014172	(HSPC150), mRNA
NM_014173	Homo sapiens HSPC142 protein (HSPC142), mRNA
NM_014172	Homo sapiens HSPC141 protein (HSPC141), mRNA
NM_014171	Homo sapiens postsynaptic protein CRIPT (CRIPT), mRNA
NM_014169	Homo sapiens HSPC134 protein (HSPC134), mRNA
NM_014168	Homo sapiens HSPC133 protein (HSPC133), mRNA
NM_014167	Homo sapiens HSPC128 protein (HSPC128), mRNA
NM_014165	Homo sapiens HSPC125 protein (HSPC125), mRNA
NM_014163	Homo sapiens HSPC073 protein (HSPC073), mRNA
NM_014162	Homo sapiens HSPC072 protein (HSPC072), mRNA
NM_014159	Homo sapiens Huntingtin interacting protein B (HYPB), mRNA
NM_014158	Homo sapiens HSPC067 protein (HSPC067), mRNA
NM_014157	Homo sapiens HSPC065 protein (HSPC065), mRNA
NM_014152	Homo sapiens HSPC054 protein (HSPC054), mRNA
NM_014151	Homo sapiens HSPC053 protein (HSPC053), mRNA
NM_014148	Homo sapiens HSPC048 protein (HSPC048), mRNA
NM_014147	Homo sapiens HSPC047 protein (HSPC047), mRNA
NM_014041	Homo sapiens signal peptidase 12kDa (SPC12), mRNA
NM_014047	Homo sapiens HSPC023 protein (HSPC023), mRNA
NM_014028	Homo sapiens HSPC019 protein (HSPC019), mRNA
NM_014026	Homo sapiens HSPC015 protein (HSPC015), mRNA
NM_015362	Homo sapiens HSPC002 protein (HSPC002), mRNA
NM_015603	Homo sapiens DKFZP586M1019 protein (DKFZP586M1019), mRNA
NM 015537	Homo sapiens DKFZP586J1624 protein (DKFZP586J1624), mRNA
NM_015584	Homo sapiens DKFZP586F1524 protein (DKFZP586F1524), mRNA
NM 015677	Homo sapiens hypothetical protein (DKFZP586F1318), mRNA
NM_015416	Homo sapiens DKFZP586A011 protein (DKFZP586A011), mRNA
NM_015513	Homo sapiens DKFZP566D213 protein (DKFZP566D213), mRNA
NM_015509	Homo sapiens DKFZP566B183 protein (DKFZP566B183), mRNA
NM_014042	Homo sapiens DKFZP564M082 protein (DKFZP564M082), mRNA
NM_015455	Homo sapiens KIAA1194 protein (KIAA1194), mRNA
NM 015601	Homo sapiens DKFZP564G092 protein (DKFZP564G092), mRNA
NM 014044	Homo sapiens DKFZP564G0222 protein (DKFZP564G0222), mRNA
NM 015658	Homo sapiens DKFZP564C186 protein (DKFZP564C186), mRNA
NM_015654	Homo sapiens DKFZP564C103 protein (DKFZP564C103), mRNA
NM_015535	Homo sapiens DKFZP564A2416 protein (DKFZP564A2416), mRNA
NM 014034	Homo sapiens DKFZP547E2110 protein (DKFZP547E2110), mRNA
NM 015607	Homo sapiens DKFZP547E1010 protein (DKFZP547E1010), mRNA
NM 015594	Homo sapiens DKFZP434O047 protein (DKFZP434O047), mRNA
NM_015492	Homo sapiens DKFZP434H132 protein (DKFZP434H132), mRNA
NM 015515	Homo sapiens type I intermediate filament cytokeratin (HAIK1), mRNA
NM 014064	Homo sapiens AD-003 protein (AD-003), mRNA
NM_014517	Homo sapiens upstream binding protein 1 (LBP-1a) (UBP1), mRNA
NM 014294	Homo sapiens translocating chain-associating membrane protein (TRAM),
	mRNA
NM 014305	Homo sapiens dTDP-D-glucose 4,6-dehydratase (TDPGD), mRNA
NM_014300	Homo sapiens signal peptidase complex (18kD) (SPC18), mRNA
NM 014419	Homo sapiens soggy-1 gene (DKKL1-pending), mRNA
NM_014445	Homo sapiens stress-associated endoplasmic reticulum protein 1; ribosome
	associated membrane protein 4 (SERP1), mRNA
NM 014329	Homo sapiens autoantigen (RCD-8), mRNA
	anomingen (ICOD-0), mil 477

NM_014504	Homo sapiens putative Rab5 GDP/GTP exchange factor homologue (RABEX5),
NM 014589	mRNA Homo sapiens phospholipas A2, group IIE (PLA2G2E), mRNA
NM 014471	Homo sapiens serine protease inhibitor, Kazal type 4 (SPINK4), mRNA
NM 014891	Homo sapiens PDGFA associated protein 1 (PDAP1), mRNA
NM 014308	Homo sapiens phosphoinositide-3-kinase, regulatory subunit, polypeptide p101
1111_014508	(P101-PI3K), mRNA
NM_014359	Homo sapiens opticin (OPTC), mRNA
NM_014515	Homo sapiens CCR4-NOT transcription complex, subunit 2 (CNOT2), mRNA
NM_014360	Homo sapiens NK-2 (Drosophila) homolog 8 (NKX2.8), mRNA
NM_014371	Homo sapiens neighbor of A-kinase anchoring protein 95 (NAKAP95), mRNA
NM_014342	Homo sapiens mitochondrial carrier homolog 2 (MTCH2), nuclear gene
	encoding mitochondrial protein, mRNA
NM_015716	Homo sapiens Misshapen/NIK-related kinase (MINK), mRNA
NM_014358	Homo sapiens C-type (calcium dependent, carbohydrate-recognition domain)
	lectin, superfamily member 9 (CLECSF9), mRNA
NM_014552	Homo sapiens LBP protein 32 (LBP-32), mRNA
NM_014247	Homo sapiens PDZ domain containing guanine nucleotide exchange
	factor(GEF)1 (PDZ-GEF1), mRNA
NM_014267	Homo sapiens small acidic protein (IMAGE145052), mRNA
NM_014597	Homo sapiens acidic 82 kDa protein mRNA (HSU15552), mRNA
NM_014254	Homo sapiens transmembrane protein 5 (TMEM5), mRNA
NM_014362	Homo sapiens 3-hydroxyisobutyryl-Coenzyme A hydrolase (HIBCH), mRNA
NM_014365	Homo sapiens protein kinase H11 (H11), mRNA
NM_014584	Homo sapiens ERO1-like (S. cerevisiae) (ERO1L), mRNA
NM_014367	Homo sapiens hypothetical protein, estradiol-induced (E2IG5), mRNA
NM_014366	Homo sapiens putative nucleotide binding protein, estradiol-induced (E2IG3), mRNA
NM_014380	Homo sapiens nerve growth factor receptor (TNFRSF16) associated protein 1 (NGFRAP1), mRNA
NM_014890	Homo sapiens downregulated in ovarian cancer 1 (DOC1), mRNA
NM_014595	Homo sapiens 5' nucleotidase, deoxy (pyrimidine), cytosolic type C (NT5C), mRNA
NM_014316	Homo sapiens calcium-regulated heat-stable protein (24kD) (CRHSP-24), mRNA
NM 014430	Homo sapiens cell death-inducing DFFA-like effector b (CIDEB), mRNA
NM_014400	Homo sapiens GPI-anchored metastasis-associated protein homolog (C4.4A), mRNA
NM_014408	Homo sapiens similar to yeast BET3 (S. cerevisiae) (BET3), mRNA
NM_014374	Homo sapiens replication initiation region protein (60kD) (RIP60), mRNA
NM_013943	Homo sapiens chloride intracellular channel 4 (CLIC4), mRNA
NM_013433	Homo sapiens karyopherin beta 2b, transportin (TRN2), mRNA
NM_013435	Homo sapiens retinal homeobox protein (RX), mRNA
NM 013377	Homo sapiens hypothetical protein (DKFZp434B0417), mRNA
NM 012297	Homo sapiens Ras-GTPase activating protein SH3 domain-binding protein 2
	(KIAA0660), mRNA
NM_013286	Homo sapiens chromosome 3p21.1 gene sequence (HUMAGCGB), mRNA
NM_012472	Homo sapiens testis specific leucine rich repeat protein (TSLRP), mRNA
NM_012119	Homo sapiens cell cycle related kinase (CCRK), mRNA
NM_013266	Homo sapiens alpha-catenin-like protein (VR22), mRNA
NM_013346	Homo sapiens sorting nexin 12 (SNX12), mRNA
NM_013322	Homo sapiens sorting nexin 10 (SNX10), mRNA

NM 013400	Homo sapiens replication initiation region protein (60kD) (RIP60), mRNA
NM 013355	Homo sapiens protein kinase PKNbeta (pknbeta), mRNA
NM 013240	Homo sapiens putative N6-DNA-methyltransferase (N6AMT1), mRNA
NM 013364	Homo sapiens paraneoplastic cancer-testis-brain antigen (MA5), mRNA
NM 013275	Homo sapiens nasopharyngeal carcinoma susceptibility protein (LZ16), mRNA
NM 013312	Homo sapiens hook2 protein (HOOK2), mRNA
NM 013332	Homo sapiens hypoxia-inducible protein 2 (HIG2), mRNA
NM 013308	Homo sapiens platelet activating receptor homolog (H963), mRNA
NM 013394	Homo sapiens acid fibroblast growth factor-like protein (GLIO703), mRNA
NM 013329	Homo sapiens chromosome 21 open reading frame 66 (C21orf66), mRNA
NM 013333	Homo sapiens EH domain-binding mitotic phosphoprotein (EPSIN), mRNA
NM 013395	Homo sapiens proteinx0008 (AD013), mRNA
NM_012463	Homo sapiens TJ6 protein (TJ6), mRNA
NM 012461	Homo sapiens TERF1 (TRF1)-interacting nuclear factor 2 (TINF2), mRNA
NM 012245	Homo sapiens SKI-interacting protein (SNW1), mRNA
NM 012437	Homo sapiens SNARE associated protein snapin (SNAPAP), mRNA
NM_012433	Homo sapiens splicing factor 3b, subunit 1, 155kD (SF3B1), mRNA
NM 012431	Homo sapiens sema domain, immunoglobulin domain (Ig), short basic domain,
_	secreted, (semaphorin) 3E (SEMA3E), mRNA
NM_012234	Homo sapiens RING1 and YY1 binding protein (RYBP), mRNA
NM_012420	Homo sapiens retinoic acid- and interferon-inducible protein (58kD) (RI58),
	mRNA
NM_012417	Homo sapiens retinal degeneration B beta (RDGBB), mRNA
NM_012229	Homo sapiens 5'-nucleotidase (purine), cytosolic type B (NT5B), mRNA
NM_012390	Homo sapiens protein homologous to salivary proline-rich protein P-B (PBI),
	mRNA
NM_012346	Homo sapiens nucleoporin 62kD (NUP62), mRNA
NM_012339	Homo sapiens transmembrane 4 superfamily member (tetraspan NET-7) (NET-7), mRNA
NM_012338	Homo sapiens transmembrane 4 superfamily member (tetraspan NET-2) (NET-
	2), mRNA
NM_012332	Homo sapiens Mitochondrial Acyl-CoA Thioesterase (MT-ACT48), mRNA
NM_012327	Homo sapiens phosphatidylinositol glycan, class N (PIGN), mRNA
NM_012321	Homo sapiens U6 snRNA-associated Sm-like protein (LSM4), mRNA
NM_012294	Homo sapiens guanine nucleotide exchange factor for Rap1; M-Ras-regulated GEF (KIAA0277), mRNA
NM 012289	Homo sapiens Kelch-like ECH-associated protein 1 (KIAA0132), mRNA
NM_012285	Homo sapiens potassium voltage-gated channel, subfamily H (eag-related),
	member 4 (KCNH4), mRNA
NM_012267	Homo sapiens hsp70-interacting protein (HSPBP1), mRNA
NM_012266	Homo sapiens DnaJ (Hsp40) homolog, subfamily B, member 5 (DNAJB5), mRNA
NM 012260	Homo sapiens 2-hydroxyphytanoyl-CoA lyase (HPCL2), mRNA
NM_012204	Homo sapiens general transcription factor IIIC, polypeptide 4 (90kD) (GTF3C4),
	mRNA
NM_012086	Homo sapiens general transcription factor IIIC, polypeptide 3 (102kD) (GTF3C3), mRNA
NM 012155	Homo sapiens microtubule-associated protein like echinoderm EMAP (EMAP-
_	2), mRNA
NM_012123	Homo sapiens CGI-02 protein (CGI-02), mRNA
NM_012097	Homo sapiens ADP-ribosylation factor-like 5 (ARL5), mRNA
NM_005028	Homo sapiens phosphatidylinositol-4-phosphate 5-kinase, type II, alpha
	Propries 5 Kinds, type II, uipita

	(PIP5K2A), mRNA
NM 006869	Homo sapiens centaurin, alpha 1 (CENTA1), mRNA
NM 007362	Homo sapiens nuclear cap binding protein subunit 2, 20kD (NCBP2), mRNA
NM 007358	Homo sapiens putative DNA binding protein (M96), mRNA
NM 007344	Homo sapiens transcription termination factor, RNA polymerase I (TTF1),
	mRNA
NM_007369	Homo sapiens G-protein coupled receptor (RE2), mRNA
NM_005176	Homo sapiens ATP synthase, H+ transporting, mitochondrial F0 complex, subunit c (subunit 9), isoform 2 (ATP5G2), mRNA
NM_007347	Homo sapiens adaptor-related protein complex 4, epsilon 1 subunit (AP4E1), mRNA
NM 002673	Homo sapiens plexin B1 (PLXNB1), mRNA
NM_007034	Homo sapiens DnaJ (Hsp40) homolog, subfamily B, member 4 (DNAJB4), mRNA
NM_004547	Homo sapiens NADH dehydrogenase (ubiquinone) 1 beta subcomplex, 4 (15kD, B15) (NDUFB4), mRNA
NM_007180	Homo sapiens trehalase (brush-border membrane glycoprotein) (TREH), mRNA
NM_007115	Homo sapiens tumor necrosis factor, alpha-induced protein 6 (TNFAIP6), mRNA
NM 007217	Homo sapiens programmed cell death 10 (PDCD10), mRNA
NM 007269	Homo sapiens syntaxin binding protein 3 (STXBP3), mRNA
NM 007107	Homo sapiens signal sequence receptor, gamma (translocon-associated protein
	gamma) (SSR3), mRNA
NM 007282	Homo sapiens ring finger protein 13 (RNF13), mRNA
NM_007265	Homo sapiens suppressor of S. cerevisiae gcr2 (HSGT1), mRNA
NM 007223	Homo sapiens putative G protein coupled receptor (GPR), mRNA
NM_007192	Homo sapiens chromatin-specific transcription elongation factor, 140 kDa subunit (FACTP140), mRNA
NM 007263	Homo sapiens coatomer protein complex, subunit epsilon (COPE), mRNA
NM 007005	Homo sapiens BCE-1 protein (BCE-1), mRNA
NM 007019	Homo sapiens ubiquitin-conjugating enzyme E2C (UBE2C), mRNA
NM_007064	Homo sapiens serine/threonine kinase with Dbl- and pleckstrin homology domains (TRAD), mRNA
NM_007062	Homo sapiens nuclear phosphoprotein similar to S. cerevisiae PWP1 (PWP1), mRNA
NM 007080	Homo sapiens Sm protein F (LSM6), mRNA
NM 007072	Homo sapiens HERV-H LTR-associating 2 (HHLA2), mRNA
NM_007077	Homo sapiens adaptor-related protein complex 4, sigma 1 subunit (AP4S1),
	mRNA
NM 006751	Homo sapiens sperm specific antigen 2 (SSFA2), mRNA
NM 006748	Homo sapiens Src-like-adaptor (SLA), mRNA
NM 006851	Homo sapiens glioma pathogenesis-related protein (RTVP1), mRNA
NM 006815	Homo sapiens coated vesicle membrane protein (RNP24), mRNA
NM 006741	Homo sapiens protein phosphatase 1, regulatory (inhibitor) subunit 1A
_	(PPP1R1A), mRNA
NM_006823	Homo sapiens protein kinase (cAMP-dependent, catalytic) inhibitor alpha (PKIA), mRNA
NM 006825	
	Homo sapiens cytoskeleton-associated protein 4 (CKAP4), mRNA
NM_006833	Homo sapiens COP9 subunit 6 (MOV34 homolog, 34 kD) (MOV34-34KD), mRNA
NM_006838	Homo sapiens methionyl aminopeptidase 2 (METAP2), mRNA
NM_006634	Homo sapiens vesicle-associated membrane protein 5 (myobrevin) (VAMP5),

	mRNA
NM 006676	Homo sapiens ubiquitin specific protease 20 (USP20), mRNA
NM 006662	Homo sapiens Snf2-related CBP activator protein (SRCAP), mRNA
NM 006692	Homo sapiens DNA-binding protein amplifying expression of surfactant protein
14141_000052	B (SPBPBP), mRNA
NM 006590	Homo sapiens SnRNP assembly defective 1 homolog (SAD1), mRNA
NM 006695	Homo sapiens RaP2 interacting protein 8 (RPIP8), mRNA
NM 006663	Homo sapiens RelA-associated inhibitor (RAI), mRNA
NM 006570	Homo sapiens Ras-related GTP-binding protein (RAGA), mRNA
NM 002721	Homo sapiens protein phosphatase 6, catalytic subunit (PPP6C), mRNA
NM 006627	Homo sapiens POP4 (processing of precursor, S. cerevisiae) homolog (POP4),
	mRNA
NM 006580	Homo sapiens claudin 16 (CLDN16), mRNA
NM_006648	Homo sapiens serologically defined colon cancer antigen 43 (SDCCAG43), mRNA
NM 006681	Homo sapiens neuromedin U (NMU), mRNA
NM 006554	Homo sapiens metaxin 2 (MTX2), mRNA
NM 006609	Homo sapiens mitogen-activated protein kinase kinase kinase 2 (MAP3K2),
	mRNA
NM_004274	Homo sapiens A kinase (PRKA) anchor protein 6 (AKAP6), mRNA
NM_006633	Homo sapiens IQ motif containing GTPase activating protein 2 (IQGAP2),
	mRNA
NM_006548	Homo sapiens IGF-II mRNA-binding protein 2 (IMP-2), mRNA
NM_006644	Homo sapiens heat shock 105kD (HSP105B), mRNA
NM_006543	Homo sapiens Mahlavu hepatocellular carcinoma (HHCM), mRNA
NM_006540	Homo sapiens nuclear receptor coactivator 2 (NCOA2), mRNA
NM_006578	Homo sapiens guanine nucleotide binding protein (G protein), beta 5 (GNB5), mRNA
NM_006550	Homo sapiens fibrinogen silencer binding protein (FSBP), mRNA
NM_006678	Homo sapiens CMRF35 leukocyte immunoglobulin-like receptor (CMRF35), mRNA
NM 006569	Homo sapiens cell growth regulatory with EF-hand domain (CGR11), mRNA
NM_006584	Homo sapiens chaperonin containing TCP1, subunit 6B (zeta 2) (CCT6B),
	mRNA
NM_006538	Homo sapiens BCL2-like 11 (apoptosis facilitator) (BCL2L11), mRNA
NM_006628	Homo sapiens cyclic AMP phosphoprotein, 19 kD (ARPP-19), mRNA
NM_006370	Homo sapiens vesicle-associated soluble NSF attachment protein receptor (v-SNARE; homolog of S. cerevisiae VTI1) (VTI2), mRNA
NM_006354	Homo sapiens transcriptional adaptor 3 (ADA3, yeast homolog)-like (PCAF
_	histone acetylase complex) (TADA3L), mRNA
NM_006456	Homo sapiens sialyltransferase (STHM), mRNA
NM_006409	Homo sapiens actin related protein 2/3 complex, subunit 1A (41 kD) (ARPC1A),
	mRNA
NM_006279	Homo sapiens sialyltransferase 6 (N-acetyllacosaminide alpha 2,3-
) D 6 00 55 15	sialyltransferase) (SIAT6), mRNA
NM_006142	Homo sapiens stratifin (SFN), mRNA
NM_006455	Homo sapiens nucleolar autoantigen (55kD) similar to rat synaptonemal complex
NIM OOCALA	protein (SC65), mRNA
NM_006414	Homo sapiens ribonuclease P (38kD) (RPP38), mRNA
NM 006413	Homo sapiens ribonuclease P (30kD) (RPP30), mRNA
NM 006423	Homo sapiens Rab acceptor 1 (prenylated) (RABAC1), mRNA
NM_006239	Homo sapiens protein phosphatase, EF hand calcium-binding domain 2 (PPEF2),

	mRNA
NM 006230	Homo sapiens polymerase (DNA directed), delta 2, regulatory subunit (50kD)
11212_000250	(POLD2), mRNA
NM 006156	Homo sapiens neural precursor cell expressed, developmentally down-regulated
	8 (NEDD8), mRNA
NM 006369	Homo sapiens MUF1 protein (MUF1), mRNA
NM 006441	Homo sapiens 5,10-methenyltetrahydrofolate synthetase (5-
_	formyltetrahydrofolate cyclo-ligase) (MTHFS), mRNA
NM_006309	Homo sapiens leucine rich repeat (in FLII) interacting protein 2 (LRRFIP2),
	mRNA
NM_006330	Homo sapiens lysophospholipase I (LYPLA1), mRNA
NM_006344	Homo sapiens macrophage lectin 2 (calcium dependent) (HML2), mRNA
NM_006395	Homo sapiens ubiquitin activating enzyme E1-like protein (GSA7), mRNA
NM_006322	Homo sapiens spindle pole body protein (GCP3), mRNA
NM_006141	Homo sapiens dynein, cytoplasmic, light intermediate polypeptide 2 (DNCLI2),
	mRNA
NM_006416	Homo sapiens solute carrier family 35 (CMP-sialic acid transporter), member 1
	(SLC35A1), mRNA
NM_006349	Homo sapiens putative cyclin G1 interacting protein (CG1I), mRNA
NM_006429	Homo sapiens chaperonin containing TCP1, subunit 7 (eta) (CCT7), mRNA
NM_006430	Homo sapiens chaperonin containing TCP1, subunit 4 (delta) (CCT4), mRNA
NM_006431	Homo sapiens chaperonin containing TCP1, subunit 2 (beta) (CCT2), mRNA
NM_002810	Homo sapiens proteasome (prosome, macropain) 26S subunit, non-ATPase, 4 (PSMD4), mRNA
NM_006002	Homo sapiens ubiquitin carboxyl-terminal esterase L3 (ubiquitin thiolesterase)
	(UCHL3), mRNA
NM_006068	Homo sapiens toll-like receptor 6 (TLR6), mRNA
NM_006100	Homo sapiens alpha2,3-sialyltransferase (ST3GALVI), mRNA
NM_006061	Homo sapiens specific granule protein (28 kDa) (SGP28), mRNA
NM_006063	Homo sapiens sarcomeric muscle protein (SARCOSIN), mRNA
NM_006076	Homo sapiens Rev/Rex activation domain binding protein-related (RAB-R), mRNA
NM_006034	Homo sapiens p53-induced protein (PIG11), mRNA
NM_006039	Homo sapiens endocytic receptor (macrophage mannose receptor family) (KIAA0709), mRNA
NM_006018	Homo sapiens putative chemokine receptor; GTP-binding protein (HM74), mRNA
NM_006101	Homo sapiens highly expressed in cancer, rich in leucine heptad repeats (HEC),
1111_000101	mRNA
NM_006098	Homo sapiens guanine nucleotide binding protein (G protein), beta polypeptide
_	2-like 1 (GNB2L1), mRNA
NM_005895	Homo sapiens golgi autoantigen, golgin subfamily a, 3 (GOLGA3), mRNA
NM_006023	Homo sapiens D123 gene product (D123), mRNA
NM_006090	Homo sapiens choline/ethanolaminephosphotransferase (CEPT1), mRNA
NM_005822	Homo sapiens Down syndrome critical region gene 1-like 1 (DSCR1L1), mRNA
NM_005827	Homo sapiens UDP-galactose transporter related (UGTREL1), mRNA
NM_005725	Homo sapiens tetraspan 2 (TSPAN-2), mRNA
NM_005879	Homo sapiens TRAF interacting protein (TRIP), mRNA
NM_005816	Homo sapiens T cell activation, increased late expression (TACTILE), mRNA
NM_005843	Homo sapiens signal transducing adaptor molecule (SH3 domain and ITAM motif) 2 (STAM2), mRNA
NM 005636	Homo sapiens synovial sarcoma, X breakpoint 4 (SSX4), mRNA
1.1.1 000000	1 suprems synovial salconia, A oreakpoint 4 (SSA4), HINNA

Homo sapiens vinexin beta (SH3-containing adaptor molecule-1) (SCAM-1), mRNA Homo sapiens hypothetical SBBI03 protein (SBB103), mRNA Homo sapiens stromal antigen 1 (STAG1), mRNA
Homo sapiens reticulon 2 (RTN2), mRNA
Homo sapiens ribonuclease, RNase A family, k6 (RNASE6), mRNA
Homo sapiens retinol dehydrogenase homolog (RDHL), mRNA
Homo sapiens Rab9 effector p40 (RAB9P40), mRNA
Homo sapiens phenylalanyl-tRNA synthetase beta-subunit (PheHB), mRNA
Homo sapiens protein phosphatase 3 (formerly 2B), catalytic subunit, gamma soform (calcineurin A gamma) (PPP3CC), mRNA
Homo sapiens nuclear transport factor 2 (placental protein 15) (PP15), mRNA
Homo sapiens protein disulfide isomerase-related protein (P5), mRNA
Homo sapiens 37 kDa leucine-rich repeat (LRR) protein (P37NB), mRNA
Homo sapiens STIP1 homology and U-Box containing protein 1 (STUB1), nRNA
Homo sapiens natural killer cell group 7 sequence (NKG7), mRNA
Homo sapiens nuclear domain 10 protein (NDP52), mRNA
Homo sapiens melan-A (MLANA), mRNA
Homo sapiens leucyl/cystinyl aminopeptidase (LNPEP), mRNA
Homo sapiens short-chain alcohol dehydrogenase family member (HEP27), nRNA
Homo sapiens carbohydrate (N-acetylglucosamine 6-O) sulfotransferase 4 CHST4), mRNA
Homo sapiens WD-repeat protein (HAN11), mRNA
Homo sapiens nuclear RNA helicase, DECD variant of DEAD box family DDXL), mRNA
Homo sapiens CD36 antigen (collagen type I receptor, thrombospondin eceptor)-like 1 (CD36L1), mRNA
Homo sapiens CCAAT-box-binding transcription factor (CBF2), mRNA
Homo sapiens calcitonin receptor-like (CALCRL), mRNA
Homo sapiens actin related protein 2/3 complex, subunit 1B (41 kD) (ARPC1B), nRNA
Iomo sapiens nuclear protein, marker for differentiated aortic smooth muscle nd down-regulated with vascular injury (APEG1), mRNA
Iomo sapiens heat shock 27kD protein 1 (HSPB1), mRNA
Iomo sapiens thyroid hormone receptor-associated protein, 95-kD subunit TRAP95), mRNA
Homo sapiens regulator of Fas-induced apoptosis (TOSO), mRNA
Iomo sapiens trophinin associated protein (tastin) (TROAP), mRNA
Homo sapiens signal transducer and activator of transcription 2, 113kD STAT2), mRNA
Iomo sapiens SUMO-1 activating enzyme subunit 1 (SAE1), mRNA
Iomo sapiens protein kinase C, epsilon (PRKCE), mRNA
Iomo sapiens pyruvate dehydrogenase kinase, isoenzyme 3 (PDK3), mRNA
Iomo sapiens DnaJ (Hsp40) homolog, subfamily B, member 6 (DNAJB6), nRNA
Iomo sapiens RNA polymerase II transcriptional regulation mediator (Med6, S. erevisiae, homolog of) (MED6), mRNA
lomo sapiens growth factor recentor-hound protein 7 (GRR7) mPNA
Iomo sapiens growth factor receptor-bound protein 7 (GRB7), mRNA Iomo sapiens gap junction protein, alpha 7, 45kD (connexin 45) (GJA7), mRNA

	subunit c (subunit 9), isoform 1 (ATP5G1), mRNA
NM_003418	Homo sapiens zinc finger protein 9 (a cellular retroviral nucleic acid binding
	protein) (ZNF9), mRNA
NM_005151	Homo sapiens ubiquitin specific protease 14 (tRNA-guanine transglycosylase)
	(USP14), mRNA
NM_005119	Homo sapiens thyroid hormone receptor-associated protein, 150 kDa subunit
	(TRAP150), mRNA
NM_005071	Homo sapiens solute carrier family 1 (high affinity aspartate/glutamate
	transporter), member 6 (SLC1A6), mRNA
NM_005047	Homo sapiens proteasome (prosome, macropain) 26S subunit, non-ATPase, 5
	(PSMD5), mRNA
NM_005134	Homo sapiens protein phosphatase 4, regulatory subunit 1 (PPP4R1), mRNA
NM_005033	Homo sapiens polymyositis/scleroderma autoantigen 1 (75kD) (PMSCL1),
	mRNA
NM_005025	Homo sapiens serine (or cysteine) proteinase inhibitor, clade I (neuroserpin),
	member 1 (SERPINI1), mRNA
NM_005023	Homo sapiens protein geranylgeranyltransferase type I, beta subunit (PGGT1B),
	mRNA
NM_005020	Homo sapiens phosphodiesterase 1C, calmodulin-dependent (70kD) (PDE1C),
	mRNA
NM_005017	Homo sapiens phosphate cytidylyltransferase 1, choline, alpha isoform
	(PCYT1A), mRNA
NM_005131	Homo sapiens nuclear matrix protein p84 (P84), mRNA
NM_005101	Homo sapiens interferon-stimulated protein, 15 kDa (ISG15), mRNA
NM_005122	Homo sapiens nuclear receptor subfamily 1, group I, member 3 (NR1I3), mRNA
NM_004666	Homo sapiens vanin 1 (VNN1), mRNA
NM_004247	Homo sapiens U5 snRNP-specific protein, 116 kD (U5-116KD), mRNA
NM_004704	Homo sapiens U3 snoRNP-associated 55-kDa protein (U3-55K), mRNA
NM_004786	Homo sapiens thioredoxin-like, 32kD (TXNL), mRNA
NM_004257	Homo sapiens TGF beta receptor associated protein -1 (TRAP-1), mRNA
NM_004620	Homo sapiens TNF receptor-associated factor 6 (TRAF6), mRNA
NM_004604	Homo sapiens syntaxin 4A (placental) (STX4A), mRNA
NM_004785	Homo sapiens solute carrier family 9 (sodium/hydrogen exchanger), isoform 3
	regulatory factor 2 (SLC9A3R2), mRNA
NM_004252	Homo sapiens solute carrier family 9 (sodium/hydrogen exchanger), isoform 3
	regulatory factor 1 (SLC9A3R1), mRNA
NM_004694	Homo sapiens solute carrier family 16 (monocarboxylic acid transporters),
	member 6 (SLC16A6), mRNA
NM_004696	Homo sapiens solute carrier family 16 (monocarboxylic acid transporters).
	member 4 (SLC16A4), mRNA
NM_004263	Homo sapiens sema domain, immunoglobulin domain (Ig), transmembrane
	domain (TM) and short cytoplasmic domain, (semaphorin) 4F (SEMA4F).
	mRNA
NM_004868	Homo sapiens glycoprotein, synaptic 2 (GPSN2), mRNA
NM_004844	Homo sapiens SH3-domain binding protein 5 (BTK-associated) (SH3BP5),
	mRNA
NM_004703	Homo sapiens rabaptin-5 (RAB5EP), mRNA
NM_004249	Homo sapiens RAB28, member RAS oncogene family (RAB28), mRNA
NM_004218	Homo sapiens RAB11B, member RAS oncogene family (RAB11B), mRNA
NM_004676	Homo sapiens PTPN13-like, Y-linked (PRY), mRNA
NM_004726	Homo sapiens RALBP1 associated Eps domain containing 2 (REPS2), mRNA
NM_004881	Homo sapiens quinone oxidoreductase homolog (PIG3), mRNA
	1

ND4 004671	Homo sapiens Protein inhibitor of activated STAT X (PIASX-BETA), mRNA
NM_004671	Homo sapiens peroxisomal biogenesis factor 14 (PEX14), mRNA
NM 004565	Homo sapiens peroxisomal biogenesis factor 14 (PEX14), mking Homo sapiens phosphate cytidylyltransferase 1, choline, beta isoform
NM_004845	(PCYT1B), mRNA
NM_004563	Homo sapiens phosphoenolpyruvate carboxykinase 2 (mitochondrial) (PCK2), mRNA
NM 004800	Homo sapiens transmembrane 9 superfamily member 2 (TM9SF2), mRNA
NM_004556	Homo sapiens nuclear factor of kappa light polypeptide gene enhancer in B-cells inhibitor, epsilon (NFKBIE), mRNA
NM 004647	Homo sapiens Neuro-d4 (rat) homolog (NEUD4), mRNA
NM_004546	Homo sapiens NADH dehydrogenase (ubiquinone) 1 beta subcomplex, 2 (8kD, AGGG) (NDUFB2), mRNA
NM_004545	Homo sapiens NADH dehydrogenase (ubiquinone) 1 beta subcomplex, 1 (7kD, MNLL) (NDUFB1), mRNA
NM_004542	Homo sapiens NADH dehydrogenase (ubiquinone) 1 alpha subcomplex, 3 (9kD, B9) (NDUFA3), mRNA
NM_004544	Homo sapiens NADH dehydrogenase (ubiquinone) 1 alpha subcomplex, 10 (42kD) (NDUFA10), mRNA
NM_004784	Homo sapiens N-deacetylase/N-sulfotransferase (heparan glucosaminyl) 3 (NDST3), mRNA
NM_004901	Homo sapiens lysosomal apyrase-like 1 (LYSAL1), mRNA
NM_004798	Homo sapiens kinesin family member 3B (KIF3B), mRNA
NM_004515	Homo sapiens interleukin enhancer binding factor 2, 45kD (ILF2), mRNA
NM_004838	Homo sapiens Homer, neuronal immediate early gene, 3 (HOMER-3), mRNA
NM_004854	Homo sapiens HNK-1 sulfotransferase (HNK-1ST), mRNA
NM_004488	Homo sapiens glycoprotein V (platelet) (GP5), mRNA
NM 004485	Homo sapiens guanine nucleotide binding protein 4 (GNG4), mRNA
NM 004122	Homo sapiens growth hormone secretagogue receptor (GHSR), mRNA
NM_004479	Homo sapiens fucosyltransferase 7 (alpha (1,3) fucosyltransferase) (FUT7), mRNA
NM 004438	Homo sapiens EphA4 (EPHA4), mRNA
NM_004094	Homo sapiens eukaryotic translation initiation factor 2, subunit 1 (alpha, 35kD) (EIF2S1), mRNA
NM_004681	Homo sapiens eukaryotic translation initiation factor 1A, Y chromosome (EIF1AY), mRNA
NM_004226	Homo sapiens serine/threonine kinase 17b (apoptosis-inducing) (STK17B), mRNA
NM_004792	Homo sapiens peptidyl-prolyl isomerase G (cyclophilin G) (PPIG), mRNA
NM_004831	Homo sapiens cofactor required for Sp1 transcriptional activation, subunit 7 (70kD) (CRSP7), mRNA
NM_004269	Homo sapiens cofactor required for Sp1 transcriptional activation, subunit 8 (34kD) (CRSP8), mRNA
NM_004270	Homo sapiens cofactor required for Sp1 transcriptional activation, subunit 9 (33kD) (CRSP9), mRNA
NM_004232	Homo sapiens STAT induced STAT inhibitor-4 (CIS4), mRNA
NM_004882	Homo sapiens CBF1 interacting corepressor (CIR), mRNA
NM_004198	Homo sapiens cholinergic receptor, nicotinic, alpha polypeptide 6 (CHRNA6), mRNA
NM_004825	Homo sapiens chromodomain protein, Y chromosome, 2 (CDY2), mRNA
NM_004351	Homo sapiens Cas-Br-M (murine) ectropic retroviral transforming sequence b (CBLB), mRNA
NM_004054	Homo sapiens complement component 3a receptor 1 (C3AR1), mRNA

NM_004899	Homo sapiens brain and reproductive organ-expressed (TNFRSF1A modulator) (BRE), mRNA
NM_004889	Homo sapiens ATP synthase, H+ transporting, mitochondrial F0 complex,
ND 6 00 4000	subunit f, isoform 2 (ATP5J2), mRNA
NM_004890	Homo sapiens sperm associated antigen 7 (SPAG7), mRNA
NM_004908_	Homo sapiens pre-T/NK cell associated protein (6H9A), mRNA
NM_003406	Homo sapiens tyrosine 3-monooxygenase/tryptophan 5-monooxygenase activation protein, zeta polypeptide (YWHAZ), mRNA
NM_003574	Homo sapiens VAMP (vesicle-associated membrane protein)-associated protein A (33kD) (VAPA), mRNA
NM_001073	Homo sapiens UDP glycosyltransferase 2 family, polypeptide B11 (UGT2B11), mRNA
NM 003300	Homo sapiens TNF receptor-associated factor 3 (TRAF3), mRNA
NM_003297	Homo sapiens nuclear receptor subfamily 2, group C, member 1 (NR2C1), mRNA
NM 003212	Homo sapiens teratocarcinoma-derived growth factor 1 (TDGF1), mRNA
NM 003763	Homo sapiens syntaxin 16 (STX16), mRNA
NM 003955	Homo sapiens STAT induced STAT inhibitor 3 (SSI-3), mRNA
NM 003693	Homo sapiens acetyl LDL receptor; SREC=scavenger receptor expressed by
	endothelial cells (SREC), mRNA
NM_003563	Homo sapiens speckle-type POZ protein (SPOP), mRNA
NM_003578	Homo sapiens sterol O-acyltransferase 2 (SOAT2), mRNA
NM_003099	Homo sapiens sorting nexin 1 (SNX1), mRNA
NM_003095	Homo sapiens small nuclear ribonucleoprotein polypeptide F (SNRPF), mRNA
NM_003091	Homo sapiens small nuclear ribonucleoprotein polypeptides B and B1 (SNRPB), mRNA
NM_003086	Homo sapiens small nuclear RNA activating complex, polypeptide 4, 190kD (SNAPC4), mRNA
NM_003084	Homo sapiens small nuclear RNA activating complex, polypeptide 3, 50kD (SNAPC3), mRNA
NM_003825	Homo sapiens synaptosomal-associated protein, 23kD (SNAP23), mRNA
NM_003983	Homo sapiens solute carrier family 7 (cationic amino acid transporter, y+ system), member 6 (SLC7A6), mRNA
NM_003916	Homo sapiens adaptor-related protein complex 1, sigma 2 subunit (AP1S2), mRNA
NM_003896	Homo sapiens sialyltransferase 9 (CMP-NeuAc:lactosylceramide alpha-2,3-sialyltransferase; GM3 synthase) (SIAT9), mRNA
NM 003769	Homo sapiens splicing factor, arginine/serine-rich 9 (SFRS9), mRNA
NM_003016	Homo sapiens splicing factor, arginine/serine-rich 2 (SFRS2), mRNA
NM_003161	Homo sapiens ribosomal protein S6 kinase, 70kD, polypeptide 1 (RPS6KB1), mRNA
NM_003708	Homo sapiens microsomal NAD+-dependent retinol dehydrogenase 4 (RODH-4), mRNA
NM 002933	Homo sapiens ribonuclease, RNase A family, 1 (pancreatic) (RNASE1), mRNA
NM_002919	Homo sapiens regulatory factor X, 3 (influences HLA class II expression) (RFX3), mRNA
NM_002865	Homo sapiens RAB2, member RAS oncogene family (RAB2), mRNA
NM 002849	Homo sapiens protein tyrosine phosphatase, receptor type, R (PTPRR), mRNA
NM 002822	Homo sapiens protein tyrosine kinase 9 (PTK9), mRNA
NM_002812	Homo sapiens proteasome (prosome, macropain) 26S subunit, non-ATPase, 8 (PSMD8), mRNA
NM 002808	Homo sapiens proteasome (prosome, macropain) 26S subunit, non-ATPase, 2
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NM_002816		(PSMD2), mRNA
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NM 002157 Homo sapiens heat shock 10kD protein 1 (chaperonin 10) (HSPE1), mRNA		
		Homo sapiens heat shock 10kD protein 1 (chaperonin 10) (USDE1) DNA
NM_001521 Homo sapiens general transcription factor IIIC, polypeptide 2 (beta subunit, 110kD) (GTF3C2), mRNA	NM_001521	Homo sapiens general transcription factor IIIC, polypeptide 2 (beta subunit,
	NM_001516	

	(GTF2H3), mRNA
NM 003910	Homo sapiens maternal G10 transcript (G10), mRNA
NM 001969	Homo sapiens eukaryotic translation initiation factor 5 (EIF5), mRNA
NM 003751	Homo sapiens eukaryotic translation initiation factor 3, subunit 9 (eta, 116kD)
	(EIF3S9), mRNA
NM 003755	Homo sapiens eukaryotic translation initiation factor 3, subunit 4 (delta, 44kD)
] -	(EIF3S4), mRNA
NM_003756	Homo sapiens eukaryotic translation initiation factor 3, subunit 3 (gamma, 40kD)
-	(EIF3S3), mRNA
NM 001414	Homo sapiens eukaryotic translation initiation factor 2B, subunit 1 (alpha, 26kD)
_	(EIF2B1), mRNA
NM 001412	Homo sapiens eukaryotic translation initiation factor 1A (EIF1A), mRNA
NM_003566	Homo sapiens early endosome antigen 1, 162kD (EEA1), mRNA
NM_001957	Homo sapiens endothelin receptor type A (EDNRA), mRNA
NM_001936	Homo sapiens dipeptidylpeptidase VI (DPP6), mRNA
NM_003648	Homo sapiens diacylglycerol kinase, delta (130kD) (DGKD), mRNA
NM_001921	Homo sapiens dCMP deaminase (DCTD), mRNA
NM_003590	Homo sapiens cullin 3 (CUL3), mRNA
NM_003592	Homo sapiens cullin 1 (CUL1), mRNA
NM_001207	Homo sapiens basic transcription factor 3 (BTF3), mRNA
NM_001191	Homo sapiens BCL2-like 1 (BCL2L1), mRNA
NM_001689	Homo sapiens ATP synthase, H+ transporting, mitochondrial F0 complex,
	subunit c (subunit 9) isoform 3 (ATP5G3), mRNA
NM_001688	Homo sapiens ATP synthase, H+ transporting, mitochondrial F0 complex,
	subunit b, isoform 1 (ATP5F1), mRNA
NM_003664	Homo sapiens adaptor-related protein complex 3, beta 1 subunit (AP3B1), mRNA
NM 058168	Homo sapiens gene differentially expressed in prostate (GDEP), mRNA
NM 058222	Homo sapiens tectorin beta (TECTB), mRNA
NM 058192	Homo sapiens ribosomal large subunit pseudouridine synthase C like (RLUCL),
	mRNA
NM 058190	Homo sapiens chromosome 21 open reading frame 70 (C21orf70), mRNA
NM 058189	Homo sapiens chromosome 21 open reading frame 69 (C21orf69), mRNA
NM 058186	Homo sapiens chromosome 21 open reading frame 11 (C21orf11), mRNA
NM 058184	Homo sapiens chromosome 21 open reading frame 42 (C21orf42), mRNA
NM_058182	Homo sapiens chromosome 21 open reading frame 51 (C21orf51), mRNA
NM 058180	Homo sapiens chromosome 21 open reading frame 58 (C21orf58), mRNA
NM_058173	Homo sapiens small breast epithelial mucin (LOC118430), mRNA
NM_058172	Homo sapiens capillary morphogenesis protein 2 (CMG2), mRNA
NM_017884	Homo sapiens PIN2-interacting protein 1 (PINX1), mRNA
NM_054021	Homo sapiens G protein-coupled receptor 101 (GPR101), mRNA
NM_053280	Homo sapiens h-Shippo 1 (LOC113746), mRNA
NM_003449	Homo sapiens tripartite motif-containing 26 (TRIM26), mRNA
NM_052939	Homo sapiens Fc receptor-like protein 3 (FCRH3), mRNA
NM_052938	Homo sapiens Fc receptor-like protein 1 (FCRH1), mRNA
NM_052872	Homo sapiens interleukin 17F (IL17F), mRNA
NM_024011	Homo sapiens cell division cycle 2-like 2 (CDC2L2), transcript variant 1, mRNA
NM_033621	Homo sapiens cell division cycle 2-like 2 (CDC2L2), transcript variant 10, mRNA
NM 033537	
NM 033536	Homo sapiens cell division cycle 2-like 2 (CDC2L2), transcript variant 9, mRNA
NM 033534	Homo sapiens cell division cycle 2-like 2 (CDC2L2), transcript variant 8, mRNA
141AT 022224	Homo sapiens cell division cycle 2-like 2 (CDC2L2), transcript variant 7, mRNA

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NM_033532	Homo sapiens cell division cycle 2-like 2 (CDC2L2), transcript variant 6, mRNA
NM_033531	Homo sapiens cell division cycle 2-like 2 (CDC2L2), transcript variant 5, mRNA
NM_033529	Homo sapiens cell division cycle 2-like 2 (CDC2L2), transcript variant 4, mRNA
NM_033528	Homo sapiens cell division cycle 2-like 2 (CDC2L2), transcript variant 3, mRNA
NM_033527	Homo sapiens cell division cycle 2-like 2 (CDC2L2), transcript variant 2, mRNA
NM_006629	Homo sapiens zinc finger protein 271 (ZNF271), mRNA
NM_015294	Homo sapiens tripartite motif-containing 37 (TRIM37), mRNA
NM_033132	Homo sapiens zinc family member 5 protein (ZIC5), mRNA
NM_033108	Homo sapiens heat shock transcription factor 2-like (LOC86614), mRNA
NM_033106	Homo sapiens galanin-like peptide precursor (LOC85569), mRNA
NM_033105	Homo sapiens beta cysteine string protein (LOC85479), mRNA
NM_033104	Homo sapiens stonin 2 (LOC85439), mRNA
NM_033102	Homo sapiens prostein protein (LOC85414), mRNA
NM_003823	Homo sapiens tumor necrosis factor receptor superfamily, member 6b, decoy
	(TNFRSF6B), transcript variant M68E, mRNA
NM_006470	Homo sapiens tripartite motif-containing 16 (TRIM16), mRNA
NM_032606	Homo sapiens calcyphosine (LOC84698), mRNA
NM_032595	Homo sapiens neurabin II (LOC84687), mRNA
NM_032584	Homo sapiens zinc finger protein 347 (ZNF347), mRNA
NM_032576	Homo sapiens lipopolysaccaride-specific response 5-like protein (LOC84663), mRNA
NM_032518	Homo sapiens collagen-like Alzheimer amyloid plaque component precursor (LOC84570), mRNA
NM_032509	Homo sapiens RNA binding protein (LOC84549), mRNA
NM_032484	Homo sapiens hypothetical protein (LOC84514), mRNA
NM_032389	Homo sapiens zinc finger protein 289, ID1 regulated (ZNF289), mRNA
NM_031918	Homo sapiens Kruppel-like factor 16 (KLF16), mRNA
NM_031463	Homo sapiens steroid dehydrogenase-like (LOC83693), mRNA
NM_031461	Homo sapiens CocoaCrisp (LOC83690), mRNA
NM_031417	Homo sapiens MAP/microtubule affinity-regulating kinase like 1 (MARKL1), mRNA
NM 030791	Homo sapiens sphingosine-1-phosphatase (LOC81537), mRNA
NM_024670	Homo sapiens suppressor of variegation 3-9 (Drosophila) homolog 2;
	hypothetical protein FLJ23414 (SUV39H2), mRNA
NM_003414	Homo sapiens zinc finger protein 267 (ZNF267), transcript variant 498723, mRNA
NM_023945	Homo sapiens membrane-spanning 4-domains, subfamily A, member 5 (MS4A5), mRNA
NM_023014	Homo sapiens hypothetical protein similar to preferentially expressed antigen of melanoma (LOC65122), mRNA
NM_023013	Homo sapiens hypothetical protein similar to preferentially expressed antigen of melanoma (LOC65121), mRNA
NM_022357	Homo sapiens putative metallopeptidase (family M19) (LOC64180), mRNA
NM_022355	Homo sapiens putative dipeptidase (LOC64174), mRNA
NM_022353	Homo sapiens putative sialoglycoprotease type 2 (LOC64172), mRNA
NM_022345	Homo sapiens uterine-derived 14 kDa protein (LOC64150), mRNA
NM_022343	Homo sapiens 17kD fetal brain protein (LOC64148), mRNA
NM_022340	Homo sapiens FYVE-finger-containing Rab5 effector protein rabenosyn-5
	(LOC64145), mRNA
NM_021932	Homo sapiens hypothetical protein from EUROIMAGE 1987170 (LOC60626), mRNA
NM 021931	Homo sapiens DEAD/H (Asp-Glu-Ala-Asp/His) box polypeptide 35 (DDX35),
	the contraction of polypepine 33 (DDA33);

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	mRNA
NM_021632	Homo sapiens zinc-finger protein ZBRK1 (ZBRK1), mRNA
NM_021630	Homo sapiens PDZ-LIM protein mystique (LOC59346), mRNA
NM_019591	Homo sapiens zinc finger protein 26 (KOX 20) (ZNF26), mRNA
NM_018675	Homo sapiens zinc finger protein 302 (ZNF302), mRNA
NM_021226	Homo sapiens hypothetical protein from clones 23549 and 23762 (LOC58504),
	mRNA
NM_021211	Homo sapiens transposon-derived Buster1 transposase-like protein (LOC58486),
	mRNA
NM_021186	Homo sapiens zona pellucida glycoprotein 4 (ZP4), mRNA
NM_020903	Homo sapiens ubiquitin-specific processing protease (LOC57663), mRNA
NM_020666	Homo sapiens CDC-like kinase 4 (CLK4), mRNA
NM_020421	Homo sapiens hypothetical protein (LOC57143), mRNA
NM_020140	Homo sapiens putative 47 kDa protein (LOC56899), mRNA
NM_016305	Homo sapiens synovial sarcoma translocation gene on chromosome 18-like 2 (SS18L2), mRNA
NM 016417	Homo sapiens clone FLB4739 (LOC51218), mRNA
NM_020467	Homo sapiens hypothetical protein from clone 643 (LOC57228), mRNA
NM 020389	Homo sapiens putative capacitative calcium channel (trp7), mRNA
NM_020385	Homo sapiens XPMC2 protein (LOC57109), mRNA
NM 020381	Homo sapiens candidate tumor suppressor protein (LOC57107), mRNA
NM 020372	Homo sapiens organic cation transporter (LOC57100), mRNA
NM 020158	Homo sapiens exosome component Rrp46 (RRP46), mRNA
NM_020147	Homo sapiens hypothetical protein from EUROIMAGE 511235 (LOC56906), mRNA
NM 020154	Homo sapiens chromosome 11 hypothetical protein ORF3 (LOC56851), mRNA
NM_019613	Homo sapiens hypothetical protein 628 (LOC56270), mRNA
NM 019059	Homo sapiens 6.2 kd protein (LOC54543), mRNA
NM 019037	Homo sapiens exosome component Rrp41 (FLJ20591), mRNA
NM 018579	Homo sapiens mitochondrial solute carrier (LOC51312), mRNA
NM 018485	Homo sapiens G protein-coupled receptor C5L2 (LOC55868), mRNA
NM_018479	Homo sapiens uncharacterized hypothalamus protein HCDASE (LOC55862), mRNA
NM 018447	Homo sapiens 30 kDa protein (LOC55831), mRNA
NM 018443	Homo sapiens zinc finger protein 302 (ZNF302), mRNA
NM 018430	Homo sapiens hypothetical protein (LOC55815), mRNA
NM 018402	Homo sapiens interleukin 26 (IL26), mRNA
NM_017692	Homo sapiens aprataxin (APTX), mRNA
NM 018171	Homo sapiens hypothetical protein FLJ10659 (FLJ10659), mRNA
NM_017530	Homo sapiens hypothetical protein LOC55565 (LOC55565), mRNA
NM_013385	Homo sapiens pleckstrin homology, Sec7 and coiled/coil domains 4 (PSCD4), mRNA
NM_016651	Homo sapiens heptacellular carcinoma novel gene-3 protein (LOC51339), mRNA
NM 016955	Homo sapiens soluble liver antigen/liver pancreas antigen (LOC51091), mRNA
NM 016422	Homo sapiens C3HC4-like zinc finger protein (ZFP26), mRNA
NM_016520	Homo sapiens hepatocellular carcinoma-associated antigen 59 (LOC51759), mRNA
NM_016275	Homo sapiens selenoprotein T (LOC51714), mRNA
NM 016242	Homo sapiens endomucin-2 (LOC51705), mRNA
NM 016233	Homo sapiens peptidylarginine deiminase type III (LOC51702), mRNA
NM 016209	Homo sapiens unknown (LOC51693), mRNA
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NM_016140	Homo sapiens brain specific protein (LOC51673), mRNA
NM_016107	Homo sapiens zinc finger RNA binding protein (ZFR), mRNA
NM 016098	Homo sapiens HSPC040 protein (LOC51660), mRNA
NM 016095	Homo sapiens HSPC037 protein (LOC51659), mRNA
NM_016086	Homo sapiens map kinase phosphatase-like protein MK-STYX (LOC51657),
_	mRNA
NM_016061	Homo sapiens CGI-127 protein (LOC51646), mRNA
NM_016039	Homo sapiens CGI-99 protein (LOC51637), mRNA
NM_016029	Homo sapiens CGI-86 protein (LOC51635), mRNA
NM_016024	Homo sapiens CGI-79 protein (LOC51634), mRNA
NM_016019	Homo sapiens CGI-74 protein (LOC51631), mRNA
NM_015964	Homo sapiens brain specific protein (LOC51673), mRNA
NM_015939	Homo sapiens CGI-09 protein (LOC51605), mRNA
NM_016647	Homo sapiens mesenchymal stem cell protein DSCD75 (LOC51337), mRNA
NM_016646	Homo sapiens mesenchymal stem cell protein DSCD28 (LOC51336), mRNA
NM_016632	Homo sapiens ARF protein (LOC51326), mRNA
NM_016629	Homo sapiens hypothetical protein (LOC51323), mRNA
NM_016627	Homo sapiens hypothetical protein (LOC51321), mRNA
NM_016626	Homo sapiens hypothetical protein (LOC51320), mRNA
NM_016618	Homo sapiens hypothetical protein (LOC51315), mRNA
NM_016616	Homo sapiens NM23-H8 (LOC51314), mRNA
NM_016613	Homo sapiens AD021 protein (LOC51313), mRNA
NM_016612	Homo sapiens mitochondrial solute carrier (LOC51312), mRNA
NM_016594	Homo sapiens FK506 binding protein precursor (LOC51303), mRNA
NM_016562	Homo sapiens toll-like receptor 7 (TLR7), mRNA
NM_016546	Homo sapiens complement C1r-like proteinase precursor, (LOC51279), mRNA
NM_016534	Homo sapiens apoptosis-related protein PNAS-1 (LOC51275), mRNA
NM_016521	Homo sapiens E2F-like protein (LOC51270), mRNA
NM_016511	Homo sapiens C-type lectin-like receptor-1 (LOC51267), mRNA
NM_016509	Homo sapiens C-type lectin-like receptor-2 (LOC51266), mRNA
NM_016496	Homo sapiens hypothetical protein (LOC51257), mRNA
NM_016494	Homo sapiens hypothetical protein (LOC51255), mRNA
NM_016484	Homo sapiens hypothetical protein (LOC51248), mRNA
NM_016471	Homo sapiens hypothetical protein (LOC51242), mRNA
NM_016467	Homo sapiens hypothetical protein (LOC51240), mRNA
NM_016454	Homo sapiens hypothetical protein (LOC51234), mRNA
NM_016429	Homo sapiens COPZ2 for nonclathrin coat protein zeta-COP (LOC51226), mRNA
NM_016383	Homo sapiens HOM-TES-85 tumor antigen (LOC51213), mRNA
NM_016380	Homo sapiens diferentiation-related protein dif13 (LOC51212), mRNA
NM_016364	Homo sapiens protein phosphatase (LOC51207), mRNA
NM_016339	Homo sapiens Link guanine nucleotide exchange factor II (LOC51195), mRNA
NM_016338	Homo sapiens Ran binding protein 11 (LOC51194), mRNA
NM_016331	Homo sapiens zinc finger protein ANC_2H01 (LOC51193), mRNA
NM_016311	Homo sapiens ATPase inhibitor precursor (LOC51189), mRNA
NM_016256	Homo sapiens N-acetylglucosamine-1-phosphodiester alpha-N-acetylglucosaminidase (LOC51172), mRNA
NM 016223	Homo sapiens protein kinase C and casein kinase substrate in neurons 3
1111_010225	(PACSIN3), mRNA
NM_016202	Homo sapiens LDL induced EC protein (LOC51157), mRNA
NM_016175	Homo sapiens truncated calcium binding protein (LOC51149), mRNA
NM 016162	Homo sapiens candidate tumor suppressor p33 ING1 homolog (LOC51147),

	DNA
ND4 016160	mRNA
NM_016158	Homo sapiens erythrocyte transmembrane protein (LOC51145), mRNA
NM_016142	Homo sapiens steroid dehydrogenase homolog (LOC51144), mRNA
NM_016141	Homo sapiens dynein light chain-A (LOC51143), mRNA
NM_016125	Homo sapiens PTD016 protein (LOC51136), mRNA
NM_016121	Homo sapiens NY-REN-45 antigen (LOC51133), mRNA
NM_016102	Homo sapiens tripartite motif-containing 17 (TRIM17), mRNA
NM_016038	Homo sapiens CGI-97 protein (LOC51119), mRNA
NM_016035	Homo sapiens CGI-92 protein (LOC51117), mRNA
NM_016026	Homo sapiens CGI-82 protein (LOC51109), mRNA
NM_016010	Homo sapiens CGI-62 protein (LOC51101), mRNA
NM_016001	Homo sapiens CGI-48 protein (LOC51096), mRNA
NM_015996	Homo sapiens CGI-40 protein (LOC51092), mRNA
NM_015978	Homo sapiens putative protein-tyrosine kinase (LOC51086), mRNA
NM_015962	Homo sapiens CGI-35 protein (LOC51077), mRNA
NM_015960	Homo sapiens CGI-32 protein (LOC51076), mRNA
NM_015957	Homo sapiens CGI-29 protein (LOC51074), mRNA
NM_015954	Homo sapiens CGI-26 protein (LOC51071), mRNA
NM_015917	Homo sapiens glutathione S-transferase subunit 13 homolog (LOC51064), mRNA
NM_015913	Homo sapiens hypothetical protein (LOC51060), mRNA
NM_015912	Homo sapiens hypothetical protein (LOC51059), mRNA
NM_015911	Homo sapiens hypothetical protein (LOC51058), mRNA
NM_015907	Homo sapiens leucine aminopeptidase (LOC51056), mRNA
NM_015883	Homo sapiens clone 1900 unknown protein (LOC51049), mRNA
NM_015872	Homo sapiens kruppel-related zinc finger protein hcKrox (LOC51043), mRNA
NM_015871	Homo sapiens zinc finger protein (LOC51042), mRNA
NM_016072	Homo sapiens CGI-141 protein (LOC51026), mRNA
NM_016068	Homo sapiens CGI-135 protein (LOC51024), mRNA
NM_016053	Homo sapiens CGI-116 protein (LOC51019), mRNA
NM_016046	Homo sapiens homolog of yeast exosomal core protein CSL4 (CSL4), mRNA
NM_016042	Homo sapiens exosome component Rrp40 (RRP40), mRNA
NM_015944	Homo sapiens CGI-14 protein (LOC51005), mRNA
NM_016060	Homo sapiens CGI-125 protein (LOC51003), mRNA
NM_016482	Homo sapiens hepatocellular carcinoma-associated antigen 59 (LOC51759), mRNA
NM_014681	Homo sapiens DEAD/H (Asp-Glu-Ala-Asp/His) box polypeptide 34 (DDX34), mRNA
NM 014415	Homo sapiens zinc finger protein (ZNF-U69274), mRNA
NM 014579	Homo sapiens zinc transporter (ZIP2), mRNA
NM_014347	Homo sapiens zinc finger protein (ZF5128), mRNA
NM_007146	Homo sapiens zinc finger protein 161 (ZNF161), mRNA
NM_006626	Homo sapiens zinc finger protein with interaction domain (ZID), mRNA
NM_006336	Homo sapiens ZYG homolog (ZYG), mRNA
NM_006138	Homo sapiens membrane-spanning 4-domains, subfamily A, member 3
37.6.005745	(hematopoietic cell-specific) (MS4A3), mRNA
NM_005741	Homo sapiens zinc finger protein 263 (ZNF263), mRNA
NM_000227	Homo sapiens laminin, alpha 3 (nicein (150kD), kalinin (165kD), BM600
27 6 000 400	(150kD), epilegrin) (LAMA3), mRNA
NM_000423	Homo sapiens keratin 2A (epidermal ichthyosis bullosa of Siemens) (KRT2A), mRNA
NM_000659	Homo sapiens autoimmune regulator (automimmune polyendocrinopathy

l dystrophy) (AIRE), transcript variant 3, mRNA
mune regulator (automimmune polyendocrinopathy
Il dystrophy) (AIRE), transcript variant AIRE-2, mRNA
mune regulator (automimmune polyendocrinopathy
Il dystrophy) (AIRE), transcript variant AIRE-1, mRNA
nger protein 177 (ZNF177), mRNA
nger protein 345 (ZNF345), mRNA
nger protein 36, C3H type, homolog (mouse) (ZFP36),
homolog, subunit of RNA polymerase III transcription
S.cerevisiae) (BRF1), mRNA
dase, beta; acid (includes glucosylceramidase) (GBA),
LOC117584), mRNA
ophic lateral sclerosis 2 (juvenile) chromosome region,
R19), mRNA
al pentraxin receptor (NPTXR), transcript variant 2, mRNA
al pentraxin receptor (NPTXR), transcript variant 1, mRNA
IB (MYO1B), mRNA
precursor cell expressed, developmentally down-regulated
RNA
family member 1B (KIF1B), mRNA
carrier family 9 (sodium/hydrogen exchanger), isoform 7
, (
sialophosphoprotein (DSPP), mRNA
elin converting enzyme 2 (ECE2), mRNA
musculoaponeurotic fibrosarcoma oncogene homolog B
NA
ss-type MMTV integration site family, member 4 (WNT4),
ss-type MMTV integration site family, member 5B
variant 1, mRNA
ss-type MMTV integration site family, member 5B
variant 2, mRNA
ss-type MMTV integration site family, member 5A
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ss-type MMTV integration site family, member 16
variant 1, mRNA
ss-type MMTV integration site family, member 16
variant 2, mRNA
te motif-containing 29 (TRIM29), transcript variant 1,
5 (, , , , , , , , , , , , , , ,
te motif-containing 29 (TRIM29), transcript variant 2,
,
nal protein L22 (RPL22), mRNA
II-like acid DNase (DLAD), transcript variant 2, mRNA
II-like acid DNase (DLAD), transcript variant 1, mRNA
en, type VI, alpha 2 (COL6A2), transcript variant 2C2a',
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en, type VI, alpha 2 (COL6A2), transcript variant 2C2a,

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NM 005061 Homo same encoding NM_030811 Homo same encoding NM_022497 Homo same encoding NM_053023 Homo same MNA NM_052826 Homo same MNA NM_052825 Homo same MNA NM_052821 Homo same MNA NM_052990 Homo same MNA NM_052989 Homo same MNA NM_052985 Homo same MNA NM_031902 Homo same MNA NM_015969 Homo same MNA NM_016065 Homo same MNA NM_031280 Homo same MNA NM_016034 Homo same MNA NM_016070 Homo same MNA NM_020191 Homo same MNA NM_021996 Homo same MNA NM_052815 Homo same MNA NM_003897 Homo same MNA NM_003897 Homo same MNA	piens ribosomal protein L4 (RPL4), mRNA
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NM_016065 Homo sa encoding NM_031280 Homo sa encoding NM_022839 Homo sa encoding NM_016034 Homo sa encoding NM_016070 Homo sa encoding NM_020191 Homo sa encoding NM_018135 Homo sa encoding NM_021996 Homo sa encoding NM_052815 Homo sa mRNA NM_003897 Homo sa mRNA	g mitochondrial protein, mRNA
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nM_016070 Homo sa encoding NM_020191 Homo sa encoding NM_018135 Homo sa encoding NM_021996 Homo sa nRNA NM_052815 Homo sa mRNA NM_003897 Homo sa mRNA	piens mitochondrial ribosomal protein S2 (MRPS2), nuclear gene
nM_020191 Homo sa encoding NM_018135 Homo sa encoding NM_021996 Homo sa nM_052815 Homo sa mRNA NM_003897 Homo sa mRNA	g mitochondrial protein, mRNA
NM_020191 Homo sa encoding NM_018135 Homo sa encoding NM_021996 Homo sa mRNA NM_052815 Homo sa mRNA NM_003897 Homo sa mRNA	piens mitochondrial ribosomal protein S23 (MRPS23), nuclear gene
encoding NM_018135 Homo sa encoding NM_021996 Homo sa mRNA NM_003897 Homo sa mRNA NM_0	g mitochondrial protein, mRNA
NM_018135 Homo same encoding NM_021996 Homo same same encoding NM_052815 Homo same encoding NM_052815 Homo same encoding MRNA NM_003897 Homo same encoding MRNA MRNA	piens mitochondrial ribosomal protein S22 (MRPS22), nuclear gene
encoding NM_021996 Homo sa MM_052815 Homo sa mRNA NM_003897 Homo sa mRNA	g mitochondrial protein, mRNA
NM_021996 Homo sa NM_052815 Homo sa mRNA NM_003897 Homo sa mRNA mRNA	piens mitochondrial ribosomal protein S18A (MRPS18A), nuclear gene
NM_052815 Homo sz mRNA NM_003897 Homo sz mRNA	g mitochondrial protein, mRNA
mRNA NM_003897 Homo sz mRNA	piens Forssman glycolipid synthetase (FS), mRNA
NM_003897 Homo sz mRNA	piens immediate early response 3 (IER3), transcript variant long,
mRNA	
	piens immediate early response 3 (IER3), transcript variant short,
NM 053013 Homo sa	
	piens enolase 3, (beta, muscle) (ENO3), transcript variant 2, mRNA
	piens CTD (carboxy-terminal domain, RNA polymerase II, polypeptide
NM_001976 Homo sa NM_048368 Homo sa A) phos	piens enolase 3, (beta, muscle) (ENO3), transcript variant 1, mRNA piens CTD (carboxy-terminal domain, RNA polymerase II, polypeptide phatase, subunit 1 (CTDP1), transcript variant FCP1b, mRNA

	1 1 1 1 1 1 (OTDD1) 4 1 1 TOD1 DATA
\	A) phosphatase, subunit 1 (CTDP1), transcript variant FCP1a, mRNA
NM_015719	Homo sapiens collagen, type V, alpha 3 (COL5A3), mRNA
NM_000393	Homo sapiens collagen, type V, alpha 2 (COL5A2), mRNA
NM_000093	Homo sapiens collagen, type V, alpha 1 (COL5A1), mRNA
NM_001256	Homo sapiens cell division cycle 27 (CDC27), mRNA
NM_004661	Homo sapiens CDC23 (cell division cycle 23, yeast, homolog) (CDC23), mRNA
NM_037370	Homo sapiens cyclin D-type binding-protein 1 (CCNDBP1), transcript variant 2, mRNA
NM_012142	Homo sapiens cyclin D-type binding-protein 1 (CCNDBP1), transcript variant 1, mRNA
NM_019592	Homo sapiens ring finger protein 20 (RNF20), mRNA
NM_003386	Homo sapiens zonadhesin (ZAN), mRNA
NM_001959	Homo sapiens eukaryotic translation elongation factor 1 beta 2 (EEF1B2), transcript variant 1, mRNA
NM_021121	Homo sapiens eukaryotic translation elongation factor 1 beta 2 (EEF1B2), transcript variant 2, mRNA
NM 006778	Homo sapiens ring finger protein 9 (RNF9), transcript variant 1, mRNA
NM 052828	Homo sapiens ring finger protein 9 (RNF9), transcript variant 2, mRNA
NM_007028	Homo sapiens tripartite motif-containing 31 (TRIM31), transcript variant 1, mRNA
NG 000019	Homo sapiens chorionic gonadotropin beta region (CGB@) on chromosome 19
NM 052952	Homo sapiens disrupted in renal carcinoma 1 (DIRC1), mRNA
NM 000989	Homo sapiens ribosomal protein L30 (RPL30), mRNA
NM 000978	Homo sapiens ribosomal protein L23 (RPL23), mRNA
NM 000985	Homo sapiens ribosomal protein L17 (RPL17), mRNA
NM 019035	Homo sapiens protocadherin 18 (PCDH18), mRNA
NM 017809	Homo sapiens nuclear RNA export factor 2 (NXF2), transcript variant 1, mRNA
NM 030943	Homo sapiens amnionless protein (AMN), mRNA
NM 022053	Homo sapiens nuclear RNA export factor 2 (NXF2), transcript variant 2, mRNA
NM 014762	Homo sapiens 24-dehydrocholesterol reductase (DHCR24), mRNA
NM 023922	Homo sapiens taste receptor, type 2, member 14 (TAS2R14), mRNA
NM 023921	Homo sapiens taste receptor, type 2, member 10 (TAS2R10), mRNA
NM 023920	Homo sapiens taste receptor, type 2, member 13 (TAS2R13), mRNA
NM_023919	Homo sapiens taste receptor, type 2, member 7 (TAS2R7), mRNA
NM 023918	Homo sapiens taste receptor, type 2, member 8 (TAS2R8), mRNA
NM_023917	Homo sapiens taste receptor, type 2, member 9 (TAS2R9), mRNA
NM 022100	Homo sapiens mitochondrial ribosomal protein S14 (MRPS14), nuclear gene
	encoding mitochondrial protein, mRNA
NM 022169	Homo sapiens ATP-binding cassette, sub-family G (WHITE), member 4
_	(ABCG4), mRNA
NM 018031	Homo sapiens WD repeat domain 6 (WDR6), transcript variant 1, mRNA
NM 012333	Homo sapiens c-myc binding protein (MYCBP), mRNA
NM_014586	Homo sapiens hormonally upregulated Neu-associated kinase (HUNK), mRNA
NM 014296	Homo sapiens calpain 7 (CAPN7), mRNA
NM_006615	Homo sapiens calpain 9 (nCL-4) (CAPN9), mRNA
NM_005807	Homo sapiens proteoglycan 4, (megakaryocyte stimulating factor, articular
	superficial zone protein, camptodactyly, arthropathy, coxa vara, pericarditis
	syndrome) (PRG4), mRNA
NM_004467	Homo sapiens fibrinogen-like 1 (FGL1), mRNA
NM_003391	Homo sapiens wingless-type MMTV integration site family member 2 (WNT2), mRNA
NM 002995	Homo sapiens small inducible cytokine subfamily C, member 1 (lymphotactin)
1111 002775	1 220110 Suprous Sman municiple cytokine Subtaining C, member 1 (1911) motacting

	(SCYC1), mRNA
NM 002477	Homo sapiens myosin, light polypeptide 5, regulatory (MYL5), mRNA
NM 058253	Homo sapiens ribosomal protein S6 kinase, 52kD, polypeptide 1 (RPS6KC1),
14141_036233	mRNA
NM 000623	Homo sapiens bradykinin receptor B2 (BDKRB2), mRNA
NM 000424	Homo sapiens keratin 5 (epidermolysis bullosa simplex, Dowling-
	Meara/Kobner/Weber-Cockayne types) (KRT5), mRNA
NM 002272	Homo sapiens keratin 4 (KRT4), mRNA
NM 057088	Homo sapiens keratin 3 (KRT3), mRNA
NM 006121	Homo sapiens keratin 1 (epidermolytic hyperkeratosis) (KRT1), mRNA
NM 057182	Homo sapiens cyclin E1 (CCNE1), transcript variant 2, mRNA
NM 001238	Homo sapiens cyclin E1 (CCNE1), transcript variant 1, mRNA
NM 054029	Homo sapiens chromosome 8 open reading frame 14 (C8orf14), mRNA
NM 054017	Homo sapiens chromosome 8 open reading frame 12 (C8orf12), mRNA
NM_052936	Homo sapiens AUT-like 2, cysteine endopeptidase (S. cerevisiae) (AUTL2), mRNA
NM 004926	Homo sapiens zinc finger protein 36, C3H type-like 1 (ZFP36L1), mRNA
NM 006887	Homo sapiens zinc finger protein 36, C3H type-like 2 (ZFP36L2), mRNA
NM 015355	Homo sapiens joined to JAZF1 (JJAZ1), mRNA
NM_005642	Homo sapiens TAF7 RNA polymerase II, TATA box binding protein (TBP)-
1111_000012	associated factor, 55 kD (TAF7), mRNA
NM 032685	Homo sapiens hypothetical protein MGC13005 (MGC13005), mRNA
NM_032656	Homo sapiens DEAD/H (Asp-Glu-Ala-Asp/His) box polypeptide 37 (DDX37),
	mRNA
NM_031919	Homo sapiens cystatin and DUF19 domain containing 1 (CSDUFD1), mRNA
NM_031475	Homo sapiens espin (ESPN), mRNA
NM_024101	Homo sapiens melanophilin (MLPH), mRNA
NM_002597	Homo sapiens phosducin (PDC), transcript variant Phd, mRNA
NM_021201	Homo sapiens membrane-spanning 4-domains, subfamily A, member 7 (MS4A7), mRNA
NM_020634	Homo sapiens growth differentiation factor 3 (GDF3), mRNA
NM_020185	Homo sapiens mitogen-activated protein kinase phosphatase x (MKPX), mRNA
NM_002897	Homo sapiens RNA binding motif, single stranded interacting protein 1
	(RBMS1), transcript variant scr2, mRNA
NM_016839	Homo sapiens RNA binding motif, single stranded interacting protein 1
	(RBMS1), transcript variant MSSP-2, mRNA
NM_016838	Homo sapiens RNA binding motif, single stranded interacting protein 1
	(RBMS1), transcript variant MSSP-1, mRNA
NM_016837	Homo sapiens RNA binding motif, single stranded interacting protein 1
3D 6 016006	(RBMS1), transcript variant MSSP-3, mRNA
NM_016836	Homo sapiens RNA binding motif, single stranded interacting protein 1
ND 6 016041	(RBMS1), transcript variant YC1, mRNA
NM_016941	Homo sapiens delta-like 3 (Drosophila) (DLL3), mRNA
NM_016335	Homo sapiens proline dehydrogenase (oxidase) 1 (PRODH), mRNA
NM_014122	Homo sapiens PRO0245 protein (PRO0245), mRNA
NM_015344	Homo sapiens leptin receptor overlapping transcript-like 1 (LEPROTL1), mRNA
NM_014450	Homo sapiens SHP2 interacting transmembrane adaptor (SIT), mRNA
NM_007159	Homo sapiens sarcolemma associated protein (SLMAP), mRNA
NM_005974	Homo sapiens proline dehydrogenase (oxidase) 1 (PRODH), mRNA
NM_004974	Homo sapiens potassium voltage-gated channel, shaker-related subfamily, member 2 (KCNA2), mRNA
NM_003195	Homo sapiens transcription elongation factor A (SII), 2 (TCEA2), mRNA

NM_001010	Homo sapiens ribosomal protein S6 (RPS6), mRNA
NM_000981	Homo sapiens ribosomal protein L19 (RPL19), mRNA
NM_003378	Homo sapiens VGF nerve growth factor inducible (VGF), mRNA
NM_001612	Homo sapiens acrosomal vesicle protein 1 (ACRV1), transcript variant 1, mRNA
NM_020115	Homo sapiens acrosomal vesicle protein 1 (ACRV1), transcript variant 11,
	mRNA
NM 020114	Homo sapiens acrosomal vesicle protein 1 (ACRV1), transcript variant 9, mRNA
NM 020113	Homo sapiens acrosomal vesicle protein 1 (ACRV1), transcript variant 8, mRNA
NM_020112	Homo sapiens acrosomal vesicle protein 1 (ACRV1), transcript variant 7, mRNA
NM 020111	Homo sapiens acrosomal vesicle protein 1 (ACRV1), transcript variant 6, mRNA
NM 020110	Homo sapiens acrosomal vesicle protein 1 (ACRV1), transcript variant 10,
	mRNA
NM 020109	Homo sapiens acrosomal vesicle protein 1 (ACRV1), transcript variant 5, mRNA
NM 020108	Homo sapiens acrosomal vesicle protein 1 (ACRV1), transcript variant 4, mRNA
NM 020107	Homo sapiens acrosomal vesicle protein 1 (ACRV1), transcript variant 3, mRNA
NM 020069	Homo sapiens acrosomal vesicle protein 1 (ACRV1), transcript variant 2, mRNA
NM 022909	Homo sapiens centromere protein H (CENPH), mRNA
NM 021734	Homo sapiens solute carrier family 25 (mitochondrial deoxynucleotide carrier),
-	member 19 (SLC25A19), mRNA
NM 021259	Homo sapiens transmembrane protein 8 (five membrane-spanning domains)
_	(TMEM8), mRNA
NM 020139	Homo sapiens oxidoreductase UCPA (LOC56898), mRNA
NM 015975	Homo sapiens TAF9-like RNA polymerase II, TATA box binding protein
_	(TBP)-associated factor, 31 kD (TAF9L), mRNA
NM 013271	Homo sapiens proprotein convertase subtilisin/kexin type 1 inhibitor (PCSK1N),
-	mRNA
NM_000904	Homo sapiens NAD(P)H dehydrogenase, quinone 2 (NQO2), mRNA
NM_000903	Homo sapiens NAD(P)H dehydrogenase, quinone 1 (NQO1), mRNA
NM_002959	Homo sapiens sortilin 1 (SORT1), mRNA
NM_057170	Homo sapiens G protein-coupled receptor kinase-interactor 2 (GIT2), transcript
	variant 2, mRNA
NM_057169	Homo sapiens G protein-coupled receptor kinase-interactor 2 (GIT2), transcript
	variant 1, mRNA
NM_057161	Homo sapiens testis intracellular mediator protein (PEAS), mRNA
NM_057167	Homo sapiens collagen, type VI, alpha 3 (COL6A3), transcript variant 5, mRNA
NM_057166	Homo sapiens collagen, type VI, alpha 3 (COL6A3), transcript variant 4, mRNA
NM_057165	Homo sapiens collagen, type VI, alpha 3 (COL6A3), transcript variant 3, mRNA
NM_057164	Homo sapiens collagen, type VI, alpha 3 (COL6A3), transcript variant 2, mRNA
NM_014776	Homo sapiens G protein-coupled receptor kinase-interactor 2 (GIT2), transcript
	variant 3, mRNA
NM_004369	Homo sapiens collagen, type VI, alpha 3 (COL6A3), transcript variant 1, mRNA
NM_001183	Homo sapiens ATPase, H+ transporting, lysosomal (vacuolar proton pump),
	subunit 1 (ATP6S1), mRNA
NM_000675	Homo sapiens adenosine A2a receptor (ADORA2A), mRNA
NM_033027	Homo sapiens AXIN1 up-regulated (AXUD1), mRNA
NM_002539	Homo sapiens ornithine decarboxylase 1 (ODC1), mRNA
NM_058004	Homo sapiens phosphatidylinositol 4-kinase, catalytic, alpha polypeptide
	(PIK4CA), transcript variant 2, mRNA
NM_000992	Homo sapiens ribosomal protein L29 (RPL29), mRNA
NM_000984	Homo sapiens ribosomal protein L23a (RPL23A), mRNA
NM_001289	Homo sapiens chloride intracellular channel 2 (CLIC2), mRNA
NM_018648	Homo sapiens nucleolar protein family A, member 3 (H/ACA small nucleolar

	DND-VAIOLA2) DNA
27 (071047	RNPs) (NOLA3), mRNA
NM_021947	Homo sapiens serine racemase (SRR), mRNA
NM_016579	Homo sapiens 8D6 antigen (8D6A), mRNA Homo sapiens protein disulfide isomerase, pancreatic (PDIP), mRNA
NM_006849	Homo sapiens protein distinde isomerase, pancteauc (1DH), microx Homo sapiens phosphatidylinositol 4-kinase, catalytic, alpha polypeptide
NM_002650	
	(PIK4CA), transcript variant 1, mRNA
NM_000988	Homo sapiens ribosomal protein L27 (RPL27), mRNA
NM_000987	Homo sapiens ribosomal protein L26 (RPL26), mRNA
NM_000986	Homo sapiens ribosomal protein L24 (RPL24), mRNA
NM_031964	Homo sapiens keratin associated protein 17.1 (KAP17.1), mRNA
NM_000420	Homo sapiens Kell blood group (KEL), mRNA Homo sapiens serine/threonine kinase 22C (spermiogenesis associated)
NM_052841	(STK22C), mRNA
NM_017647	Homo sapiens FtsJ homolog 3 (E. coli) (FTSJ3), mRNA
NM_001845	Homo sapiens collagen, type IV, alpha 1 (COL4A1), mRNA
NM_016508	Homo sapiens cyclin-dependent kinase-like 3 (CDKL3), mRNA
NM_001261	Homo sapiens cyclin-dependent kinase 9 (CDC2-related kinase) (CDK9), mRNA
NM_033131	Homo sapiens wingless-type MMTV integration site family, member 3A (WNT3A), mRNA
NM_030753	Homo sapiens wingless-type MMTV integration site family, member 3 (WNT3), mRNA
NM_003396	Homo sapiens wingless-type MMTV integration site family, member 15 (WNT15), mRNA
NM_004626	Homo sapiens wingless-type MMTV integration site family, member 11 (WNT11), mRNA
NM 057176	Homo sapiens barttin (BSND), mRNA
NM_012079	Homo sapiens diacylglycerol O-acyltransferase homolog 1 (mouse) (DGAT1), mRNA
NM 005490	Homo sapiens SH2 domain-containing 3A (SH2D3A), mRNA
NM_032563	Homo sapiens epidermal differentiation complex protein like protein (LEP16),
ND 6 014014	mRNA Homo sapiens centaurin, gamma 2 (CENTG2), mRNA
NM_014914	Homo sapiens mitochondrial ribosomal protein L18 (MRPL18), mRNA
NM_014161 NM_004895	Homo sapiens cold autoinflammatory syndrome 1 (CIAS1), mRNA
NM 000086	Homo sapiens ceroid-lipofuscinosis, neuronal 3, juvenile (Batten, Spielmeyer-
14141_000000	Vogt disease) (CLN3), mRNA
NM 033341	Homo sapiens baculoviral IAP repeat-containing 8 (BIRC8), mRNA
NM_054013	Homo sapiens mannosyl (alpha-1,3-)-glycoprotein beta-1,4-N-
1414_054015	acetylglucosaminyltransferase, isoenzyme B (MGAT4B), transcript variant 2, mRNA
NM_000449	Homo sapiens regulatory factor X, 5 (influences HLA class II expression) (RFX5), mRNA
NM_054025	Homo sapiens beta-1,3-glucuronyltransferase 1 (glucuronosyltransferase P)
14141_034023	(B3GAT1), transcript variant 2, mRNA
NM 002628	Homo sapiens profilin 2 (PFN2), transcript variant 2, mRNA
NM 053024	Homo sapiens profilin 2 (PFN2), transcript variant 1, mRNA
NM_003930	Homo sapiens src family associated phosphoprotein 2 (SCAP2), mRNA
NM_014018	Homo sapiens mitochondrial ribosomal protein S28 (MRPS28), nuclear gene
	encoding mitochondrial protein, mRNA
NM_015971	Homo sapiens mitochondrial ribosomal protein S7 (MRPS7), nuclear gene encoding mitochondrial protein, mRNA
NM 032476	Homo sapiens mitochondrial ribosomal protein S6 (MRPS6), nuclear gene
14141 032410	1 Aromo supreme introduction intospinar protein so (and so), Marie 8

· · · · · · · · · · · · · · · · · · ·	encoding mitochondrial protein, mRNA
NM 018141	Homo sapiens mitochondrial ribosomal protein S10 (MRPS10), nuclear gene
11112_010111	encoding mitochondrial protein, mRNA
NM 014046	Homo sapiens mitochondrial ribosomal protein S18B (MRPS18B), nuclear gene
1111_011010	encoding mitochondrial protein, mRNA
NM 006513	Homo sapiens seryl-tRNA synthetase (SARS), mRNA
NM 021153	Homo sapiens cadherin 19, type 2 (CDH19), mRNA
NM 033664	Homo sapiens cadherin 11, type 2, OB-cadherin (osteoblast) (CDH11), transcript
14141_033004	variant 2, mRNA
NM_001797	Homo sapiens cadherin 11, type 2, OB-cadherin (osteoblast) (CDH11), transcript
	variant 1, mRNA
NM_033381	Homo sapiens collagen, type IV, alpha 5 (Alport syndrome) (COL4A5),
_	transcript variant 3, mRNA
NM 033380	Homo sapiens collagen, type IV, alpha 5 (Alport syndrome) (COL4A5),
_	transcript variant 2, mRNA
NM_000495	Homo sapiens collagen, type IV, alpha 5 (Alport syndrome) (COL4A5),
_	transcript variant 1, mRNA
NM_000092	Homo sapiens collagen, type IV, alpha 4 (COL4A4), mRNA
NM 033184	Homo sapiens keratin associated protein 2.4 (KAP2.4), mRNA
NM_032014	Homo sapiens mitochondrial ribosomal protein S24 (MRPS24), nuclear gene
_	encoding mitochondrial protein, mRNA
NM 001006	Homo sapiens ribosomal protein S3A (RPS3A), mRNA
NM_012411	Homo sapiens protein tyrosine phosphatase, non-receptor type 22 (lymphoid)
-	(PTPN22), transcript variant 2, mRNA
NM 015967	Homo sapiens protein tyrosine phosphatase, non-receptor type 22 (lymphoid)
-	(PTPN22), transcript variant 1, mRNA
NM 006310	Homo sapiens aminopeptidase puromycin sensitive (NPEPPS), mRNA
NM_033335	Homo sapiens nuclear receptor subfamily 6, group A, member 1 (NR6A1),
-	transcript variant 3, mRNA
NM_033334	Homo sapiens nuclear receptor subfamily 6, group A, member 1 (NR6A1),
-	transcript variant 1, mRNA
NM_001489	Homo sapiens nuclear receptor subfamily 6, group A, member 1 (NR6A1),
_	transcript variant 2, mRNA
NM_001606	Homo sapiens ATP-binding cassette, sub-family A (ABC1), member 2
_	(ABCA2), mRNA
NM 002284	Homo sapiens keratin, hair, basic, 6 (monilethrix) (KRTHB6), mRNA
NM 002283	Homo sapiens keratin, hair, basic, 5 (KRTHB5), mRNA
NM 002282	Homo sapiens keratin, hair, basic, 3 (KRTHB3), mRNA
NM 033033	Homo sapiens keratin, hair, basic, 2 (KRTHB2), mRNA
NM 002281	Homo sapiens keratin, hair, basic, 1 (KRTHB1), mRNA
NM 033045	Homo sapiens keratin, hair, basic, 4 (KRTHB4), mRNA
NM_001011	Homo sapiens ribosomal protein S7 (RPS7), mRNA
NM 000980	Homo sapiens ribosomal protein L18a (RPL18A), mRNA
NM_000979	Homo sapiens ribosomal protein L18 (RPL18), mRNA
NM 000977	Homo sapiens ribosomal protein L13 (RPL13), transcript variant 1, mRNA
NM 033251	Homo sapiens ribosomal protein L13 (RPL13), transcript variant 2, mRNA
NM 000976	Homo sapiens ribosomal protein L12 (RPL12), mRNA
NM 000975	Homo sapiens ribosomal protein L12 (RI L12), mRNA Homo sapiens ribosomal protein L11 (RPL11), mRNA
NM 000973	Homo sapiens hossomar protein ETT (RF ETT), mixtx Homo sapiens luteinizing hormone beta polypeptide (LHB), mRNA
NM 005082	Homo sapiens zinc finger protein 147 (estrogen-responsive finger protein)
_	(ZNF147), mRNA
NM_003549	Homo sapiens hyaluronoglucosaminidase 3 (HYAL3), mRNA

NIM 022191	Homo sapiens cannabinoid receptor 1 (brain) (CNR1), transcript variant 3,
NM_033181	mRNA
NG_000018	Homo sapiens genomic type I (acidic) hair keratin gene cluster (KRTHA.1@) on
	chromosome 17
NM 033151	Homo sapiens ATP-binding cassette, sub-family C (CFTR/MRP), member 11
	(ABCC11), mRNA
NM_006998	Homo sapiens secretagogin (SECRET), mRNA
NM_006201	Homo sapiens PCTAIRE protein kinase 1 (PCTK1), transcript variant 1, mRNA
NM_033019	Homo sapiens PCTAIRE protein kinase 1 (PCTK1), transcript variant 3, mRNA
NM_033018	Homo sapiens PCTAIRE protein kinase 1 (PCTK1), transcript variant 2, mRNA
NG_000012	Homo sapiens genomic protocadherin gamma cluster (PCDHG@) on
_	chromosome 5
NM_001023	Homo sapiens ribosomal protein S20 (RPS20), mRNA
NM_004451	Homo sapiens estrogen-related receptor alpha (ESRRA), mRNA
NM_005755	Homo sapiens Epstein-Barr virus induced gene 3 (EBI3), mRNA
NM_001015	Homo sapiens ribosomal protein S11 (RPS11), mRNA
NM_006923	Homo sapiens stromal cell-derived factor 2 (SDF2), mRNA
NM_000394	Homo sapiens crystallin, alpha A (CRYAA), mRNA
NM_003761	Homo sapiens vesicle-associated membrane protein 8 (endobrevin) (VAMP8),
	mRNA
NM_031958	Homo sapiens keratin associated protein 3.1 (KRTAP3.1), mRNA
NM_031957	Homo sapiens keratin associated protein 1.5 (KRTAP1.5), mRNA
NM_004776	Homo sapiens UDP-Gal:betaGlcNAc beta 1,4- galactosyltransferase, polypeptide
	5 (B4GALT5), mRNA
NM_030587	Homo sapiens UDP-Gal:betaGlcNAc beta 1,4- galactosyltransferase, polypeptide
	2 (B4GALT2), transcript variant 1, mRNA
NM_003780	Homo sapiens UDP-Gal:betaGlcNAc beta 1,4- galactosyltransferase, polypeptide
37.6 00 100	2 (B4GALT2), transcript variant 2, mRNA
NM_004391	Homo sapiens cytochrome P450, subfamily VIIIB (sterol 12-alpha-hydroxylase),
ND4 000705	polypeptide 1 (CYP8B1), mRNA
NM_000785	Homo sapiens cytochrome P450, subfamily XXVIIB (25-hydroxyvitamin D-1-
	alpha-hydroxylase), polypeptide 1 (CYP27B1), mitochondrial protein encoded
NM_031419	by nuclear gene, mRNA Homo sapiens molecule possessing ankyrin repeats induced by
14147 021413	lipopolysaccharide (MAIL), homolog of mouse (MAIL), mRNA
NM 000961	Homo sapiens prostaglandin I2 (prostacyclin) synthase (PTGIS), mRNA
NM 003293	Homo sapiens tryptase, alpha (TPS1), mRNA
NM_016630	Homo sapiens acid cluster protein 33 (ACP33), mRNA
NM 014458	Homo sapiens Kelch motif containing protein (AB026190), mRNA
NM_007207	Homo sapiens dual specificity phosphatase 10 (DUSP10), mRNA
NM 030660	Homo sapiens Machado-Joseph disease (spinocerebellar ataxia 3,
	olivopontocerebellar ataxia 3, autosomal dominant, ataxia 3) (MJD), transcript
	variant 2, mRNA
NM_022055	Homo sapiens potassium channel, subfamily K, member 12 (KCNK12), mRNA
NM 021175	Homo sapiens hepcidin antimicrobial peptide (HAMP), mRNA
NM_018666	Homo sapiens sarcoma antigen (SAGE), mRNA
NM_016532	Homo sapiens SKIP for skeletal muscle and kidney enriched inositol
	phosphatase (LOC51763), mRNA
NM_015987	Homo sapiens heme binding protein 1 (HEBP1), mRNA
NM 014079	Homo sapiens Kruppel-like factor 15 (KLF15), mRNA
NM_014759	Homo sapiens phytanoyl-CoA hydroxylase interacting protein (PHYHIP),
	mRNA
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NM_002590	Homo sapiens protocadherin 8 (PCDH8), transcript variant 1, mRNA
NM_004826	Homo sapiens endothelin converting enzyme-like 1 (ECEL1), mRNA
NM_004420	Homo sapiens dual specificity phosphatase 8 (DUSP8), mRNA
NM_001012	Homo sapiens ribosomal protein S8 (RPS8), mRNA
NM_002595	Homo sapiens PCTAIRE protein kinase 2 (PCTK2), mRNA
NM_001395	Homo sapiens dual specificity phosphatase 9 (DUSP9), mRNA
NM_003887	Homo sapiens development and differentiation enhancing factor 2 (DDEF2), mRNA
NM_001446	Homo sapiens fatty acid binding protein 7, brain (FABP7), mRNA
NM_001259	Homo sapiens cyclin-dependent kinase 6 (CDK6), mRNA
NM_001760	Homo sapiens cyclin D3 (CCND3), mRNA
NM_001759	Homo sapiens cyclin D2 (CCND2), mRNA
NM_001237	Homo sapiens cyclin A2 (CCNA2), mRNA
NM_057158	Homo sapiens dual specificity phosphatase 4 (DUSP4) transcript variant 2, mRNA
NM_001394	Homo sapiens dual specificity phosphatase 4 (DUSP4), transcript variant 1, mRNA
NM_052988	Homo sapiens cyclin-dependent kinase (CDC2-like) 10 (CDK10), transcript variant 3, mRNA
NM_052987	Homo sapiens cyclin-dependent kinase (CDC2-like) 10 (CDK10), transcript variant 2, mRNA
NM_057160	Homo sapiens artemin (ARTN), transcript variant 3, mRNA
NM_057091	Homo sapiens artemin (ARTN), transcript variant 2, mRNA
NM_057090	Homo sapiens artemin (ARTN), transcript variant 4, mRNA
NM_003976	Homo sapiens artemin (ARTN), transcript variant 1, mRNA
NM_000050	Homo sapiens argininosuccinate synthetase (ASS), transcript variant 1; mRNA
NM_054012	Homo sapiens argininosuccinate synthetase (ASS), transcript variant 2, mRNA
NM_053286	Homo sapiens aquaporin 6, kidney specific (AQP6), transcript variant 2, mRNA
NM_001652	Homo sapiens aquaporin 6, kidney specific (AQP6), transcript variant 1, mRNA
NM_053032	Homo sapiens myosin, light polypeptide kinase (MYLK), transcript variant 8, mRNA
NM_053031	Homo sapiens myosin, light polypeptide kinase (MYLK), transcript variant 7, mRNA
NM_053030	Homo sapiens myosin, light polypeptide kinase (MYLK), transcript variant 5, mRNA
NM_053029	Homo sapiens myosin, light polypeptide kinase (MYLK), transcript variant 4, mRNA
NM_053028	Homo sapiens myosin, light polypeptide kinase (MYLK), transcript variant 3B, mRNA
NM_053027	Homo sapiens myosin, light polypeptide kinase (MYLK), transcript variant 3A, mRNA
NM_053026	Homo sapiens myosin, light polypeptide kinase (MYLK), transcript variant 2, mRNA
NM_053025	Homo sapiens myosin, light polypeptide kinase (MYLK), transcript variant 1, mRNA
NM_016497	Homo sapiens mitochondrial ribosomal protein 64 (MRP64), nuclear gene encoding mitochondrial protein, mRNA
NM 024026	Homo sapiens mitochondrial ribosomal protein 63 (MRP63), nuclear gene
	encoding mitochondrial protein, mRNA
NM_021821	Homo sapiens mitochondrial ribosomal protein S35 (MRPS35), nuclear gene encoding mitochondrial protein, mRNA
NM 005965	Homo sapiens myosin, light polypeptide kinase (MYLK), transcript variant 6,
1.1.1	o-passio myoom, ngm posypopuso amaso (wit i exe), mansompt variant 0,

	mRNA
NM 016640	Homo sapiens mitochondrial ribosomal protein S30 (MRPS30), mRNA
NM_053035	Homo sapiens mitochondrial ribosomal protein S33 (MRPS33), transcript variant
1411_055055	2, nuclear gene encoding mitochondrial protein, mRNA
NM 016071	Homo sapiens mitochondrial ribosomal protein S33 (MRPS33), transcript variant
1111_0100/1	1, nuclear gene encoding mitochondrial protein, mRNA
NM_031901	Homo sapiens mitochondrial ribosomal protein S21 (MRPS21), transcript variant
	1, nuclear gene encoding mitochondrial protein, mRNA
NM_018997	Homo sapiens mitochondrial ribosomal protein S21 (MRPS21), transcript variant
_	2, nuclear gene encoding mitochondrial protein, mRNA
NM 033363	Homo sapiens mitochondrial ribosomal protein S12 (MRPS12), transcript variant
_	3, nuclear gene encoding mitochondrial protein, mRNA
NM 033362	Homo sapiens mitochondrial ribosomal protein S12 (MRPS12), transcript variant
_	2, nuclear gene encoding mitochondrial protein, mRNA
NM_021144	Homo sapiens PC4 and SFRS1 interacting protein 1 (PSIP1), mRNA
NM_052953	Homo sapiens hypothetical protein LRP15 (LRP15), mRNA
NM_033207	Homo sapiens transmembrane protein HTMP10 (HTMP10), mRNA
NM_030649	Homo sapiens centaurin, beta 5 (CENTB5), mRNA
NM_023936	Homo sapiens mitochondrial ribosomal protein S34 (MRPS34), nuclear gene
	encoding mitochondrial protein, mRNA
NM_021107	Homo sapiens mitochondrial ribosomal protein S12 (MRPS12), transcript variant
	1, nuclear gene encoding mitochondrial protein, mRNA
NM_014322	Homo sapiens opsin 3 (encephalopsin, panopsin) (OPN3), mRNA
NM_001260	Homo sapiens cyclin-dependent kinase 8 (CDK8), mRNA
NM_003674	Homo sapiens cyclin-dependent kinase (CDC2-like) 10 (CDK10), transcript
NM 057094	variant 1, mRNA
NM_057093	Homo sapiens crystallin, beta A2 (CRYBA2), transcript variant 3, mRNA
NM_052984	Homo sapiens crystallin, beta A2 (CRYBA2), transcript variant 2, mRNA Homo sapiens cyclin-dependent kinase 4 (CDK4), transcript variant 2, mRNA
NM 000075	Homo sapiens cyclin-dependent kinase 4 (CDK4), transcript variant 2, mRNA Homo sapiens cyclin-dependent kinase 4 (CDK4), transcript variant 1, mRNA
NM 052827	Homo sapiens cyclin-dependent kinase 2 (CDK2), transcript variant 1, mRNA
NM 001798	Homo sapiens cyclin-dependent kinase 2 (CDK2), transcript variant 2, mRNA
NM 006522	Homo sapiens wingless-type MMTV integration site family, member 6 (WNT6),
1111_000322	mRNA
NM_005430	Homo sapiens wingless-type MMTV integration site family, member 1 (WNT1),
	mRNA
NM 003394	Homo sapiens wingless-type MMTV integration site family, member 10B
-	(WNT10B), mRNA
NM_025216	Homo sapiens wingless-type MMTV integration site family, member 10A
	(WNT10A), mRNA
NM_005370	Homo sapiens mel transforming oncogene (derived from cell line NK14)-RAB8
	homolog (MEL), mRNA
NM_033100	Homo sapiens MT-protocadherin (KIAA1775), mRNA
NM_005086	Homo sapiens sarcospan (Kras oncogene-associated gene) (SSPN), mRNA
NM_003737	Homo sapiens protocadherin 16 (PCDH16), mRNA
NM_018153	Homo sapiens tumor endothelial marker 8 (TEM8), transcript variant 3, mRNA
NM_053034	Homo sapiens tumor endothelial marker 8 (TEM8), transcript variant 2, mRNA
NM_005929	Homo sapiens antigen p97 (melanoma associated) identified by monoclonal
AD F CORSES	antibodies 133.2 and 96.5 (MFI2), transcript variant 1, mRNA
NM_033316	Homo sapiens antigen p97 (melanoma associated) identified by monoclonal
NTM 6 001000	antibodies 133.2 and 96.5 (MFI2), transcript variant 2, mRNA
NM_001002	Homo sapiens ribosomal protein, large, PO (RPLPO), transcript variant 1, mRNA

NM 053275	Homo sapiens ribosomal protein, large, P0 (RPLP0), transcript variant 2, mRNA
NM 054034	Homo sapiens fibronectin 1 (FN1), transcript variant 2, mRNA
NM 002026	Homo sapiens fibronectin 1 (FN1), transcript variant 1, mRNA
NM 004460	Homo sapiens fibroblast activation protein, alpha (FAP), mRNA
NM 000783	Homo sapiens cytochrome P450, subfamily XXVIA, polypeptide 1 (CYP26A1),
_	transcript variant 1, mRNA
NM 057157	Homo sapiens cytochrome P450, subfamily XXVIA, polypeptide 1 (CYP26A1),
	transcript variant 2, mRNA
NM_032211	Homo sapiens lysyl oxidase-like 4 (LOXL4), mRNA
NM_003395	Homo sapiens wingless-type MMTV integration site family, member 14
	(WNT14), mRNA
NM_033101	Homo sapiens lectin, galactoside-binding, soluble, 12 (galectin 12) (LGALS12),
	mRNA
NM_032611	Homo sapiens protein tyrosine phosphatase type IVA, member 3 (PTP4A3),
	transcript variant 1, mRNA
NM_007079	Homo sapiens protein tyrosine phosphatase type IVA, member 3 (PTP4A3),
	transcript variant 2, mRNA
NM_032208	Homo sapiens tumor endothelial marker 8 (TEM8), transcript variant 1, mRNA
NM_014644	Homo sapiens phosphodiesterase 4D interacting protein (myomegalin)
3 2 2 2 2 2 2 2 2 2 2 2 2 2 2 2 2 2 2 2	(PDE4DIP), mRNA
NM_006551	Homo sapiens lipophilin B (uteroglobin family member), prostatein-like
27.5.010000	(LPHB), mRNA
NM_012280	Homo sapiens FtsJ homolog 1 (E. coli) (FTSJ1), mRNA
NM_005209	Homo sapiens crystallin, beta A2 (CRYBA2), transcript variant 1, mRNA Homo sapiens opioid growth factor receptor (OGFR), mRNA
NM_007346	Homo sapiens opioid growth factor receptor (OGFR), micha Homo sapiens lipophilin A (uteroglobin family member) (LPHA), mRNA
NM_006552	Homo sapiens cell death-regulatory protein GRIM19 (GRIM19), mRNA
NM_015965	Homo sapiens mannosyl (alpha-1,3-)-glycoprotein beta-1,4-N-
NM_014275	acetylglucosaminyltransferase, isoenzyme B (MGAT4B), transcript variant 1,
	mRNA
NM_001872	Homo sapiens carboxypeptidase B2 (plasma, carboxypeptidase U) (CPB2),
14141_001872	transcript variant 1, mRNA
NM_016413	Homo sapiens carboxypeptidase B2 (plasma, carboxypeptidase U) (CPB2),
14141_010+15	transcript variant 2, mRNA
NM_004632	Homo sapiens death associated protein 3 (DAP3), transcript variant 2, nuclear
1117_00 1052	gene encoding mitochondrial protein, mRNA
NM_033657	Homo sapiens death associated protein 3 (DAP3), transcript variant 1, nuclear
	gene encoding mitochondrial protein, mRNA
NM 001266	Homo sapiens carboxylesterase 1 (monocyte/macrophage serine esterase 1)
_	(CES1), mRNA
NM 004287	Homo sapiens golgi SNAP receptor complex member 2 (GOSR2), transcript
	variant A, mRNA
NM_054022	Homo sapiens golgi SNAP receptor complex member 2 (GOSR2), transcript
l	variant B, mRNA
NM_002906	Homo sapiens radixin (RDX), mRNA
NM_001004	Homo sapiens ribosomal protein, large P2 (RPLP2), mRNA
NM_001003	Homo sapiens ribosomal protein, large, P1 (RPLP1), mRNA
NM_018644	Homo sapiens beta-1,3-glucuronyltransferase 1 (glucuronosyltransferase P)
	(B3GAT1), transcript variant 1, mRNA
NM_022145	Homo sapiens leucine zipper protein FKSG14 (FKSG14), mRNA
NM_013363	Homo sapiens procollagen C-endopeptidase enhancer 2 (PCOLCE2), mRNA
NM_033119	Homo sapiens naked cuticle homolog 1 (Drosophila) (NKD1), mRNA

NM_020439	Homo sapiens calcium/calmodulin-dependent protein kinase IG (CAMK1G), mRNA
NM_032158	Homo sapiens NOL1R2 protein (NOL1R2), mRNA
NM 022470	Homo sapiens p53 target zinc finger protein (WIG1), mRNA
NM 018044	Homo sapiens NOL1R protein (NOL1R), mRNA
NM 016262	Homo sapiens epsilon-tubulin (LOC51175), mRNA
NM 014239	Homo sapiens eukaryotic translation initiation factor 2B, subunit 2 (beta, 39kD)
	(EIF2B2), mRNA
NM_002308	Homo sapiens lectin, galactoside-binding, soluble, 9 (galectin 9) (LGALS9),
NR 6 000507	transcript variant short, mRNA
NM_009587	Homo sapiens lectin, galactoside-binding, soluble, 9 (galectin 9) (LGALS9),
NB4 001107	transcript variant long, mRNA
NM_001187	Homo sapiens B melanoma antigen (BAGE), mRNA
NM_022162	Homo sapiens caspase recruitment domain family, member 15 (CARD15), mRNA
NM_014733	Homo sapiens endosome-associated FYVE-domain protein (ENDOFIN), mRNA
NM_013393	Homo sapiens FtsJ homolog 2 (E. coli) (FTSJ2), mRNA
NM_006440	Homo sapiens thioredoxin reductase beta (TR), mRNA
NM_005863	Homo sapiens neuroepithelial cell transforming gene 1 (NET1), mRNA
NM_002119	Homo sapiens major histocompatibility complex, class II, DO alpha (HLA-
	DOA), mRNA
NM_021908	Homo sapiens suppression of tumorigenicity 7 (ST7), transcript variant b,
	mRNA
NM_018412	Homo sapiens suppression of tumorigenicity 7 (ST7), transcript variant a, mRNA
NM_054020	Homo sapiens putative ion channel protein CATSPER2 (CATSPER2), mRNA
NM_053281	Homo sapiens dachshund homolog 2 (Drosophila) (DACH2), mRNA
NM_031439	Homo sapiens SOX7 transcription factor (SOX7), mRNA
NM_030796	Homo sapiens hypothetical protein DKFZp564K0822 (DKFZP564K0822), mRNA
NM_025117	Homo sapiens hypothetical protein FLJ11871 (FLJ11871), mRNA
NM 014893	Homo sapiens KIAA0951 protein (KIAA0951), mRNA
NM_000113	Homo sapiens dystonia 1, torsion (autosomal dominant; torsin A) (DYT1), mRNA
NM_053055	Homo sapiens C-terminal modulator protein (CTMP), mRNA
NM 021212	Homo sapiens HCF-binding transcription factor Zhangfei (ZF), mRNA
NM_007237	Homo sapiens SP140 nuclear body protein (SP140), mRNA
NM 006368	Homo sapiens cAMP responsive element binding protein 3 (luman) (CREB3),
	mRNA
NM_005759	Homo sapiens abl-interactor 12 (SH3-containing protein) (AIP-1), mRNA
NM_052966	Homo sapiens chromosome 1 open reading frame 24 (C1orf24), mRNA
NM_013247	Homo sapiens protease, serine, 25 (PRSS25), mRNA
NM_003017	Homo sapiens splicing factor, arginine/serine-rich 3 (SFRS3), mRNA
NM_006289	Homo sapiens talin 1 (TLN1), mRNA
NM_000970	Homo sapiens ribosomal protein L6 (RPL6), mRNA
NM_003973	Homo sapiens ribosomal protein L14 (RPL14), mRNA
NM_001361	Homo sapiens dihydroorotate dehydrogenase (DHODH), nuclear gene encoding mitochondrial protein, mRNA
NM_021248	Homo sapiens cadherin-like 22 (CDH22), mRNA
NM 033224	Homo sapiens purine-rich element binding protein B (PURB), mRNA
NM 005859	Homo sapiens purine-rich element binding protein A (PURA), mRNA
NM 005022	Homo sapiens profilin 1 (PFN1), mRNA
NM 017481	Homo sapiens ubiquilin 3 (UBQLN3), mRNA
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NM_013444	Homo sapiens ubiquilin 2 (UBQLN2), mRNA
NM_053067	Homo sapiens ubiquilin 1 (UBQLN1), transcript variant 2, mRNA
NM_013438	Homo sapiens ubiquilin 1 (UBQLN1), transcript variant 1, mRNA
NM_032115	Homo sapiens potassium channel, subfamily K, member 16 (KCNK16), mRNA
NM_053284	Homo sapiens WAP, FS, Ig, KU, and NTR-containing protein (WFIKKN),
_	mRNA
NM 053278	Homo sapiens G protein-coupled receptor 102 (GPR102), mRNA
NM 053276	Homo sapiens vitrin (VIT), mRNA
NM 032649	Homo sapiens glutamate carboxypeptidase-like protein 2 (CPGL2), mRNA
NM 053012	Homo sapiens hypothetical protein (LOC114137), mRNA
NM 003268	Homo sapiens toll-like receptor 5 (TLR5), mRNA
NM 053005	Homo sapiens HCCA2 protein (HCCA2), mRNA
NM 052889	Homo sapiens CARD only protein (COP), mRNA
NM 024740	Homo sapiens disrupted in bipolar disorder 1 (DIBD1), mRNA
NM 015721	Homo sapiens gem (nuclear organelle) associated protein 4 (GEMIN4), mRNA
NM 003730	Homo sapiens ribonuclease 6 precursor (RNASE6PL), mRNA
NM 030916	Homo sapiens Ig superfamily receptor LNIR (LNIR), mRNA
NM_022358	Homo sapiens potassium channel, subfamily K, member 15 (TASK-5)
14141_022330	(KCNK15), mRNA
NM 022576	Homo sapiens phosducin (PDC), transcript variant PhLOP1, mRNA
NM 018269	Homo sapiens SIPL protein (SIPL), mRNA
	Homo sapiens spastic paraplegia 3A (autosomal dominant) (SPG3A), mRNA
NM_015915	Homo sapiens G protein-coupled receptor 74 (GPR74), mRNA
NM_053036	Homo sapiens paralemmin 2 (PALM2), mRNA
NM_053016	Homo sapiens hypothetical protein (LOC114138), mRNA
NM 053057	Homo sapiens septin 1 (SEPT1), mRNA
NM_052838	Homo sapiens solute carrier family 4, sodium bicarbonate transporter-like,
NM_032034	member 11 (SLC4A11), mRNA
NM 031899	Homo sapiens golgi phosphoprotein 5 (GOLPH5), mRNA
NM 018448	Homo sapiens TBP-interacting protein (TIP120A), mRNA
NM_016952	Homo sapiens cell adhesion molecule-related/down-regulated by oncogenes
	(CDON), mRNA
NM 053050	Homo sapiens mitochondrial ribosomal protein L53 (MRPL53), mRNA
NM 053045	Homo sapiens hypothetical protein MGC14327 (MGC14327), mRNA
NM 017680	Homo sapiens asporin (LRR class 1) (ASPN), mRNA
NM 003914	Homo sapiens cyclin A1 (CCNA1), mRNA
NM 032387	Homo sapiens protein kinase, lysine deficient 4 (PRKWNK4), mRNA
NM_019093	Homo sapiens UDP glycosyltransferase 1 family, polypeptide A3 (UGT1A3),
1441_017073	mRNA
NM 021027	Homo sapiens UDP glycosyltransferase 1 family, polypeptide A9 (UGT1A9),
14141_021027	mRNA
NM 019076	Homo sapiens UDP glycosyltransferase 1 family, polypeptide A8 (UGT1A8),
11112_017070	mRNA
NM 000463	Homo sapiens UDP glycosyltransferase 1 family, polypeptide A1 (UGT1A1),
11111_000703	mRNA
NM 016608	Homo sapiens ALEX1 protein (ALEX1), mRNA
NM_016607	Homo sapiens ALEX3 protein (ALEX3), mRNA
	Homo sapiens SPTF-associated factor 65 gamma (STAF65(gamma)), mRNA
NM 014860	Homo sapiens armadillo repeat protein ALEX2 (ALEX2), mRNA
NM_014782	Homo sapiens UDP glycosyltransferase 1 family, polypeptide A6 (UGT1A6),
NM_001072	mRNA
NM_000405	Homo sapiens GM2 ganglioside activator protein (GM2A), mRNA
14141 000403	Troute sapiens Out Eganghoside activator process (Ottober), and the

NM_001975	Homo sapiens enolase 2, (gamma, neuronal) (ENO2), mRNA
NM 001428	Homo sapiens enolase 1, (alpha) (ENO1), mRNA
NM 052836	Homo sapiens cadherin related 23 (CDH23), transcript variant 2, mRNA
NM 022124	Homo sapiens cadherin related 23 (CDH23), transcript variant 1, mRNA
NM 004063	Homo sapiens cadherin 17, LI cadherin (liver-intestine) (CDH17), mRNA
NM 004062	Homo sapiens cadherin 16, KSP-cadherin (CDH16), mRNA
NM 004933	Homo sapiens cadherin 15, M-cadherin (myotubule) (CDH15), mRNA
NM 001257	Homo sapiens cadherin 13, H-cadherin (heart) (CDH13), mRNA
NM_052819	Homo sapiens caspase recruitment domain protein 14 (CARD14), transcript variant 2, mRNA
NM_024110	Homo sapiens caspase recruitment domain protein 14 (CARD14), transcript variant 1, mRNA
NM_032415	Homo sapiens caspase recruitment domain family, member 11 (CARD11), mRNA
NM 014466	Homo sapiens tektin 2 (testicular) (TEKT2), mRNA
NM_053006	Homo sapiens serine/threonine kinase 22B (spermiogenesis associated) (STK22B), mRNA
NM_012083	Homo sapiens frequently rearranged in advanced T-cell lymphomas 2 (FRAT2), mRNA
NM_006922	Homo sapiens sodium channel, voltage-gated, type III, alpha polypeptide (SCN3A), mRNA
NM_005347	Homo sapiens heat shock 70kD protein 5 (glucose-regulated protein, 78kD) (HSPA5), mRNA
NM 003777	Homo sapiens dynein, axonemal, heavy polypeptide 11 (DNAH11), mRNA
NM_013282	Homo sapiens ubiquitin-like, containing PHD and RING finger domains, 1 (UHRF1), mRNA
NM 020886	Homo sapiens ubiquitin specific protease 28 (USP28), mRNA
NM_020843	Homo sapiens zinc finger protein 291 (ZNF291), mRNA
NM_024529	Homo sapiens chromosome 1 open reading frame 28 (Clorf28), mRNA
NM_053003	Homo sapiens SIGLEC-like 1 (SIGLECL1), mRNA
NM_033329	Homo sapiens SIGLEC-like 1 (SIGLECL1), mRNA
NM_015101	Homo sapiens chromosome 1 open reading frame 17 (Clorf17), mRNA
NM_032551	Homo sapiens G protein-coupled receptor 54 (GPR54), mRNA
NM_031898	Homo sapiens tektin 3 (TEKT3), mRNA
NM_025191	Homo sapiens chromosome 1 open reading frame 22 (Clorf22), mRNA
NM_022755	Homo sapiens chromosome 9 open reading frame 12 (C9orf12), mRNA
NM_021104	Homo sapiens ribosomal protein L41 (RPL41), mRNA
NM 017847	Homo sapiens chromosome 1 open reading frame 27 (Clorf27), mRNA
NM_017673	Homo sapiens chromosome 1 open reading frame 26 (Clorf26), mRNA
NM_016000	Homo sapiens mitochondrial CCA-adding tRNA-nucleotidyltransferase (MtCCA), mRNA
NM_015989	Homo sapiens cysteine sulfinic acid decarboxylase-related protein 2 (CSAD), mRNA
NM 014654	Homo sapiens syndecan 3 (N-syndecan) (SDC3), mRNA
NM 014837	Homo sapiens chromosome 1 open reading frame 16 (Clorf16), mRNA
NM 007179	Homo sapiens insulin-like 6 (INSL6), mRNA
NM 005478	Homo sapiens insulin-like 5 (INSL5), mRNA
NM 053000	Homo sapiens TIGA1 (TIGA1), mRNA
NM 052940	Homo sapiens hypothetical protein MGC8974 (MGC8974), mRNA
NM 052830	Homo sapiens gamma-glutamyltransferase-like 3 (GGTL3), mRNA
NM 053002	Homo sapiens no opposite paired repeat protein (NOPAR), mRNA
NM 052998	Homo sapiens ornithine decarboxylase-like protein (ODC-p), mRNA
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NM_052996	Homo sapiens PR domain containing 7 (PRDM7), mRNA
NM_052995	Homo sapiens Usher syndrome 3A (USH3A), mRNA
NM_007110	Homo sapiens telomerase-associated protein 1 (TEP1), mRNA
NM_033656	Homo sapiens WD repeat domain 9 (WDR9), transcript variant 2, mRNA
NM_018963	Homo sapiens WD repeat domain 9 (WDR9), transcript variant 1, mRNA
NM_017818	Homo sapiens WD repeat domain 8 (WDR8), mRNA
NM_033662	Homo sapiens WD repeat domain 4 (WDR4), transcript variant 3, mRNA
NM_033661	Homo sapiens WD repeat domain 4 (WDR4), transcript variant 2, mRNA
NM_018669	Homo sapiens WD repeat domain 4 (WDR4), transcript variant 1, mRNA
NM_017883	Homo sapiens WD repeat domain 13 (WDR13), mRNA
NM_052837	Homo sapiens secretory carrier membrane protein 3 (SCAMP3), transcript
375 005500	variant 2, mRNA
NM_005698	Homo sapiens secretory carrier membrane protein 3 (SCAMP3), transcript
\	variant 1, mRNA
NM_005697	Homo sapiens secretory carrier membrane protein 2 (SCAMP2), mRNA
NM_004866	Homo sapiens secretory carrier membrane protein 1 (SCAMP1), transcript
ND COCCCC	variant 1, mRNA
NM_052822	Homo sapiens secretory carrier membrane protein 1 (SCAMP1), transcript
ND 6 050011	variant 2, mRNA
NM_052811	Homo sapiens ret finger protein 2 (RFP2), transcript variant 2, mRNA
NM_005798	Homo sapiens ret finger protein 2 (RFP2), transcript variant 1, mRNA
NM_052817	Homo sapiens midline 2 (MID2), transcript variant 2, mRNA
NM_012216	Homo sapiens midline 2 (MID2), transcript variant 1, mRNA
NM_000798	Homo sapiens dopamine receptor D5 (DRD5), mRNA
NM_000794	Homo sapiens dopamine receptor D1 (DRD1), mRNA
NM_000796	Homo sapiens dopamine receptor D3 (DRD3), transcript variant a, mRNA
NM_033663	Homo sapiens dopamine receptor D3 (DRD3), transcript variant e, mRNA
NM_033660	Homo sapiens dopamine receptor D3 (DRD3), transcript variant d, mRNA
NM_033659	Homo sapiens dopamine receptor D3 (DRD3), transcript variant c, mRNA
NM_033658	Homo sapiens dopamine receptor D3 (DRD3), transcript variant b, mRNA
NM_004934	Homo sapiens cadherin 18, type 2 (CDH18), mRNA
NM_004061	Homo sapiens cadherin 12, type 2 (N-cadherin 2) (CDH12), mRNA
NM_030622	Homo sapiens cytochrome P450, subfamily IIS, polypeptide 1 (CYP2S1),
275 05000	mRNA
NM_052831	Homo sapiens dJ55C23.6 gene (dJ55C23.6), mRNA
NM_052816	Homo sapiens tripartite motif-containing 31 (TRIM31), transcript variant 2,
)D. C.	mRNA
NM_052812	Homo sapiens tripartite motif-containing 15 (TRIM15), transcript variant 2,
ND 6 052055	mRNA
NM_052955	Homo sapiens transglutaminase Z (TGM7), mRNA
NM_052957	Homo sapiens putative nuclear protein (NAAR1), mRNA
NM_052851	Homo sapiens similar to RhoGAP (GT650), mRNA
NM_033229	Homo sapiens tripartite motif-containing 15 (TRIM15), transcript variant 1, mRNA
NM_018103	Homo sapiens leucine-rich repeat-containing 5 (LRRC5), mRNA
NM 014879	Homo sapiens G protein-coupled receptor 105 (GPR105), mRNA
NM 000797	Homo sapiens dopamine receptor D4 (DRD4), mRNA
NM 006596	Homo sapiens polymerase (DNA directed), theta (POLQ), mRNA
NM_052972	Homo sapiens leucine-rich alpha-2-glycoprotein (LRG), mRNA
NM_052967	Homo sapiens mas-related G protein-coupled MRG (MRG), mRNA
NM_052963	Homo sapiens mitochondrial topoisomerase I (TOP1MT), mRNA
NM 052962	Homo sapiens class II cytokine receptor (IL22RA2), mRNA

NM 052954 Home sapiens vestibule-1 protein (VEST1), mRNA NM 052954 Home sapiens vestibule-1 protein (VEST1), mRNA NM 052934 Home sapiens solute carrier family 26, member 9 (SLC26A9), mRNA NM 052934 Home sapiens solute carrier family 26, member 9 (SLC26A9), mRNA NM 052934 Home sapiens proteins (RASGRP4), mRNA NM 052934 Home sapiens pro-oncosis receptor inducing membrane injury gene (PORIMIN), mRNA NM 052931 Home sapiens pro-oncosis receptor inducing membrane injury gene (PORIMIN), mRNA NM 052881 Home sapiens pro-incosis receptor inducing membrane injury gene (PORIMIN), mRNA NM 052888 Home sapiens pro-incosis receptor inducing membrane injury gene (PORIMIN), mRNA NM 052888 Home sapiens Sulta (Samana (Sa	NM_052961	Homo sapiens solute carrier family 26, member 8 (SLC26A8), mRNA
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NM_001847 Homo sapiens collagen, type IV, alpha 6 (COL4A6), transcript variant A, mRNA NM_004359 Homo sapiens cell division cycle 34 (CDC34), mRNA NM_033493 Homo sapiens cell division cycle 2-like 1 (PITSLRE proteins) (CDC2L1), transcript variant 9, mRNA NM_033492 Homo sapiens cell division cycle 2-like 1 (PITSLRE proteins) (CDC2L1), transcript variant 8, mRNA NM_033491 Homo sapiens cell division cycle 2-like 1 (PITSLRE proteins) (CDC2L1), transcript variant 7, mRNA NM_033490 Homo sapiens cell division cycle 2-like 1 (PITSLRE proteins) (CDC2L1), transcript variant 6, mRNA NM_033489 Homo sapiens cell division cycle 2-like 1 (PITSLRE proteins) (CDC2L1), transcript variant 5, mRNA NM_033488 Homo sapiens cell division cycle 2-like 1 (PITSLRE proteins) (CDC2L1), transcript variant 4, mRNA NM_033487 Homo sapiens cell division cycle 2-like 1 (PITSLRE proteins) (CDC2L1), transcript variant 3, mRNA	NM_006750	component 2) (SNTB2), mRNA
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NM_033493 Homo sapiens cell division cycle 2-like 1 (PITSLRE proteins) (CDC2L1), transcript variant 9, mRNA NM_033492 Homo sapiens cell division cycle 2-like 1 (PITSLRE proteins) (CDC2L1), transcript variant 8, mRNA NM_033491 Homo sapiens cell division cycle 2-like 1 (PITSLRE proteins) (CDC2L1), transcript variant 7, mRNA NM_033490 Homo sapiens cell division cycle 2-like 1 (PITSLRE proteins) (CDC2L1), transcript variant 6, mRNA NM_033489 Homo sapiens cell division cycle 2-like 1 (PITSLRE proteins) (CDC2L1), transcript variant 5, mRNA NM_033488 Homo sapiens cell division cycle 2-like 1 (PITSLRE proteins) (CDC2L1), transcript variant 4, mRNA NM_033487 Homo sapiens cell division cycle 2-like 1 (PITSLRE proteins) (CDC2L1), transcript variant 3, mRNA	NM_001847	
transcript variant 9, mRNA NM_033492 Homo sapiens cell division cycle 2-like 1 (PITSLRE proteins) (CDC2L1), transcript variant 8, mRNA NM_033491 Homo sapiens cell division cycle 2-like 1 (PITSLRE proteins) (CDC2L1), transcript variant 7, mRNA NM_033490 Homo sapiens cell division cycle 2-like 1 (PITSLRE proteins) (CDC2L1), transcript variant 6, mRNA NM_033489 Homo sapiens cell division cycle 2-like 1 (PITSLRE proteins) (CDC2L1), transcript variant 5, mRNA NM_033488 Homo sapiens cell division cycle 2-like 1 (PITSLRE proteins) (CDC2L1), transcript variant 4, mRNA NM_033487 Homo sapiens cell division cycle 2-like 1 (PITSLRE proteins) (CDC2L1), transcript variant 3, mRNA		
NM_033492 Homo sapiens cell division cycle 2-like 1 (PITSLRE proteins) (CDC2L1), transcript variant 8, mRNA NM_033491 Homo sapiens cell division cycle 2-like 1 (PITSLRE proteins) (CDC2L1), transcript variant 7, mRNA NM_033490 Homo sapiens cell division cycle 2-like 1 (PITSLRE proteins) (CDC2L1), transcript variant 6, mRNA NM_033489 Homo sapiens cell division cycle 2-like 1 (PITSLRE proteins) (CDC2L1), transcript variant 5, mRNA NM_033488 Homo sapiens cell division cycle 2-like 1 (PITSLRE proteins) (CDC2L1), transcript variant 4, mRNA NM_033487 Homo sapiens cell division cycle 2-like 1 (PITSLRE proteins) (CDC2L1), transcript variant 3, mRNA	NM_033493	
NM_033491 Homo sapiens cell division cycle 2-like 1 (PITSLRE proteins) (CDC2L1), transcript variant 7, mRNA NM_033490 Homo sapiens cell division cycle 2-like 1 (PITSLRE proteins) (CDC2L1), transcript variant 6, mRNA NM_033489 Homo sapiens cell division cycle 2-like 1 (PITSLRE proteins) (CDC2L1), transcript variant 5, mRNA NM_033488 Homo sapiens cell division cycle 2-like 1 (PITSLRE proteins) (CDC2L1), transcript variant 4, mRNA NM_033487 Homo sapiens cell division cycle 2-like 1 (PITSLRE proteins) (CDC2L1), transcript variant 3, mRNA	NM_033492	Homo sapiens cell division cycle 2-like 1 (PITSLRE proteins) (CDC2L1),
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NM_033489 Homo sapiens cell division cycle 2-like 1 (PITSLRE proteins) (CDC2L1), transcript variant 5, mRNA NM_033488 Homo sapiens cell division cycle 2-like 1 (PITSLRE proteins) (CDC2L1), transcript variant 4, mRNA NM_033487 Homo sapiens cell division cycle 2-like 1 (PITSLRE proteins) (CDC2L1), transcript variant 3, mRNA	NM_033490	Homo sapiens cell division cycle 2-like 1 (PITSLRE proteins) (CDC2L1),
transcript variant 5, mRNA NM_033488 Homo sapiens cell division cycle 2-like 1 (PITSLRE proteins) (CDC2L1), transcript variant 4, mRNA NM_033487 Homo sapiens cell division cycle 2-like 1 (PITSLRE proteins) (CDC2L1), transcript variant 3, mRNA	NIM 022490	
transcript variant 4, mRNA NM_033487 Homo sapiens cell division cycle 2-like 1 (PITSLRE proteins) (CDC2L1), transcript variant 3, mRNA		
NM_033487 Homo sapiens cell division cycle 2-like 1 (PITSLRE proteins) (CDC2L1), transcript variant 3, mRNA	NM_033488	
	NM_033487	Homo sapiens cell division cycle 2-like 1 (PITSLRE proteins) (CDC2L1),
	NM_033486	

	Language 1 mpNA
ND 4 001707	transcript variant 2, mRNA
NM_001787	Homo sapiens cell division cycle 2-like 1 (PITSLRE proteins) (CDC2L1),
	transcript variant 1, mRNA
NM_005983	Homo sapiens S-phase kinase-associated protein 2 (p45) (SKP2), transcript
	variant 1, mRNA
NM_032637	Homo sapiens S-phase kinase-associated protein 2 (p45) (SKP2), transcript
	variant 2, mRNA
NM_021968	Homo sapiens H4 histone family, member E (H4FE), mRNA
NM_002748	Homo sapiens mitogen-activated protein kinase 6 (MAPK6), mRNA
NM_003527	Homo sapiens H2B histone family, member N (H2BFN), mRNA
NM_001000	Homo sapiens ribosomal protein L39 (RPL39), mRNA
NM 000999	Homo sapiens ribosomal protein L38 (RPL38), mRNA
NM 000998	Homo sapiens ribosomal protein L37a (RPL37A), mRNA
NM 000997	Homo sapiens ribosomal protein L37 (RPL37), mRNA
NM 022054	Homo sapiens potassium channel, subfamily K, member 13 (KCNK13), mRNA
NM 021161	Homo sapiens potassium channel, subfamily K, member 10 (TREK-2)
	(KCNK10), mRNA
NM 003944	Homo sapiens selenium binding protein 1 (SELENBP1), mRNA
NM 033649	Homo sapiens fibroblast growth factor 18 (FGF18), transcript variant 2, mRNA
NM 004114	Homo sapiens fibroblast growth factor 13 (FGF13), transcript variant 1A, mRNA
NM 033642	Homo sapiens fibroblast growth factor 13 (FGF13), transcript variant 1B, mRNA
NM 016279	Homo sapiens cadherin 9, type 2 (T1-cadherin) (CDH9), mRNA
NM 001796	Homo sapiens cadherin 8, type 2 (CDH8), mRNA
NM 031891	Homo sapiens cadherin 20, type 2 (CDH20), mRNA
NM 006727	Homo sapiens cadherin 10, type 2 (T2-cadherin) (CDH10), mRNA
NM 033671	Homo sapiens cyclin B3 (CCNB3), transcript variant 2, mRNA
NM 033670	Homo sapiens cyclin B3 (CCNB3), transcript variant 1, mRNA
NM 033379	Homo sapiens cell division cycle 2, G1 to S and G2 to M (CDC2), transcript
14141_055577	variant 2, mRNA
NM 001786	Homo sapiens cell division cycle 2, G1 to S and G2 to M (CDC2), transcript
1111_001100	variant 1, mRNA
NM 004361	Homo sapiens cadherin 7, type 2 (CDH7), transcript variant b, mRNA
NM 033646	Homo sapiens cadherin 7, type 2 (CDH7), transcript variant a, mRNA
NM 017734	Homo sapiens palmdelphin (PALMD), mRNA
NM 052832	Homo sapiens solute carrier family 26, member 7 (SLC26A7), mRNA
NM_018718	Homo sapiens testis specific, 14 (TSGA14), mRNA
NM 015935	Homo sapiens CGI-01 protein (CGI-01), mRNA
NM 033120	Homo sapiens corot protein (CGP-07), midvA Homo sapiens naked cuticle homolog 2 (Drosophila) (NKD2), mRNA
NM 033031	Homo sapiens cyclin B3 (CCNB3), transcript variant 3, mRNA
NM 012068	Homo sapiens cyclin by (CCNB9), trained by training transcription factor 5 (ATF5), mRNA
NM 019617	
NM_019617 NM_018398	Homo sapiens CA11 (LOC56287), mRNA Homo sapiens calcium channel, voltage-dependent, alpha 2/delta 3 subunit
IMM_019339	
NM_018319	(CACNA2D3), mRNA Homo genione traced DNA phodphodiostarese (TDD1), mPNA
	Homo sapiens tyrosyl-DNA phodphodiesterase (TDP1), mRNA
NM_014404	Homo sapiens calcium channel, voltage-dependent, gamma subunit 5
NTM 014405	(CACNG5), mRNA
NM_014405	Homo sapiens calcium channel, voltage-dependent, gamma subunit 4
3D4 012114	(CACNG4), mRNA
NM_012114	Homo sapiens caspase 14, apoptosis-related cysteine protease (CASP14), mRNA
NM_006985	Homo sapiens nuclear pore complex interacting protein (NPIP), mRNA
NM_006816	Homo sapiens chromosome 5 open reading frame 8 (C5orf8), mRNA
NM_006539	Homo sapiens calcium channel, voltage-dependent, gamma subunit 3

	(CAONICE) PAIA
ND 6 004247	(CACNG3), mRNA
NM_004347	Homo sapiens caspase 5, apoptosis-related cysteine protease (CASP5), mRNA
NM_003862	Homo sapiens fibroblast growth factor 18 (FGF18), transcript variant 1, mRNA
NM_020770	Homo sapiens cingulin (KIAA1319), mRNA
NM_030778	Homo sapiens hypothetical protein PRO1331 (PRO1331), mRNA
NM_004927	Homo sapiens mitochondrial ribosomal protein L49 (MRPL49), mRNA
NM_031962	Homo sapiens keratin associated protein 9.3 (KRTAP9.3), mRNA
NM_031961	Homo sapiens keratin associated protein 9.2 (KRTAP9.2), mRNA
NM_033456	Homo sapiens potassium channel, subfamily K, member 7 (KCNK7), transcript variant E, mRNA
NM_031854	Homo sapiens keratin associated protein 4.12 (KRTAP4.12), mRNA
NM_033455	Homo sapiens potassium channel, subfamily K, member 7 (KCNK7), transcript variant D, mRNA
NM_033348	Homo sapiens potassium channel, subfamily K, member 7 (KCNK7), transcript variant B, mRNA
NM 033347	Homo sapiens potassium channel, subfamily K, member 7 (KCNK7), transcript
	variant A, mRNA
NM_033191	Homo sapiens keratin associated protein 9.4 (KAP9.4), mRNA
NM_033061	Homo sapiens keratin associated protein 4.7 (KAP4.7), mRNA
NM_033188	Homo sapiens keratin associated protein 4.5 (KAP4.5), mRNA
NM 033062	Homo sapiens keratin associated protein 4.2 (KAP4.2), mRNA
NM 033059	Homo sapiens keratin associated protein 4.14 (KAP4.14), mRNA
NM 033060	Homo sapiens keratin associated protein 4.10 (KAP4.10), mRNA
NM 033643	Homo sapiens ribosomal protein L36 (RPL36), transcript variant 1, mRNA
NM 015414	Homo sapiens ribosomal protein L36 (RPL36), transcript variant 2, mRNA
NM 007209	Homo sapiens ribosomal protein L35 (RPL35), mRNA
NM_000996	Homo sapiens ribosomal protein L35a (RPL35A), mRNA
NM 033637	Homo sapiens beta-transducin repeat containing (BTRC), transcript variant 1,
_	mRNA
NM 033345	Homo sapiens regulator of G-protein signalling 8 (RGS8), mRNA
NM 033543	Homo sapiens hypothetical protein R29124_1 (R29124_1), mRNA
NM 033186	Homo sapiens keratin associated protein 4.13 (KAP4.13), mRNA
NM 033050	Homo sapiens G protein-coupled receptor 91 (GPR91), mRNA
NM 032728	Homo sapiens hypothetical protein MGC12921 (MGC12921), mRNA
NM_032910	Homo sapiens hypothetical protein MGC14136 (MGC14136), mRNA
NM_032857	Homo sapiens mitochondrial ribosomal protein L56 (MRPL56), mRNA
NM 032640	Homo sapiens hypothetical protein MGC10526 (MGC10526), mRNA
NM_032560	Homo sapiens MSTP033 protein (MSTP033), mRNA
NM 032524	Homo sapiens keratin associated protein 4.4 (KRTAP4.4), mRNA
NM 032351	Homo sapiens mitochondrial ribosomal protein L45 (MRPL45), mRNA
NM 031963	Homo sapiens keratin associated protein 9.8 (KRTAP9.8), mRNA
NM 031432	Homo sapiens uridine-cytidine kinase 1 (UCK1), mRNA
NM 031289	Homo sapiens hypothetical protein MGC3146 (MGC3146), mRNA
NM 031269	Homo sapiens PRO1386 protein (PRO1386), mRNA
NM_030975	Homo sapiens keratin associated protein 9.9 (KRTAP9.9), mRNA
NM_030817	Homo sapiens hypothetical protein DKFZp434F0318 (DKFZP434F0318),
	mRNA
NM_017970	Homo sapiens hypothetical protein FLJ10008 (FLJ10008), mRNA
NM_024510	Homo sapiens hypothetical protein MGC4368 (MGC4368), mRNA
NM_024325	Homo sapiens hypothetical protein MGC10715 (MGC10715), mRNA
NM_023914	Homo sapiens G protein-coupled receptor 86 (GPR86), mRNA
NM_022915	Homo sapiens mitochondrial ribosomal protein L44 (MRPL44), mRNA

CHRNA10),
A.
NA
A
A
acting
, transcript
, transcript
A8), mRNA
-
NA
regulator of
_
TPase, 11
(CDK5R1),
variant 2,

encoding

NM_033640	Homo sapiens SCAN domain-containing 2 (SCAND2), transcript variant 6, mRNA
NM_033636	Homo sapiens SCAN domain-containing 2 (SCAND2), transcript variant 5, mRNA
NM_033635	Homo sapiens SCAN domain-containing 2 (SCAND2), transcript variant 4, mRNA
NM_033634	Homo sapiens SCAN domain-containing 2 (SCAND2), transcript variant 3, mRNA
NM_033633	Homo sapiens SCAN domain-containing 2 (SCAND2), transcript variant 2, mRNA
NM_022050	Homo sapiens SCAN domain-containing 2 (SCAND2), transcript variant 1, mRNA
NM 033467	Homo sapiens membrane metallo-endopeptidase-like 2 (MMEL2), mRNA
NM_032409	Homo sapiens PTEN induced putative kinase 1 (PINK1), mRNA
NM_013267	Homo sapiens breast cell glutaminase (GA), mRNA
NM_004729	Homo sapiens Ac-like transposable element (ALTE), mRNA
NM 004192	Homo sapiens acetylserotonin O-methyltransferase-like (ASMTL), mRNA
NM_002115	Homo sapiens hexokinase 3 (white cell) (HK3), nuclear gene encoding mitochondrial protein, mRNA
NM_000188	Homo sapiens hexokinase 1 (HK1), transcript variant 1, nuclear gene encoding mitochondrial protein, mRNA
NM_004728	Homo sapiens DEAD/H (Asp-Glu-Ala-Asp/His) box polypeptide 21 (DDX21), mRNA
NM_022148	Homo sapiens cytokine receptor-like factor 2 (CRLF2), mRNA
NM_022337	Homo sapiens RAB38, member RAS oncogene family (RAB38), mRNA
NM_016428	Homo sapiens NESH protein (NESH), mRNA
NM_016227	Homo sapiens chromosome 1 open reading frame 9 (C1orf9), mRNA
NM_014283	Homo sapiens chromosome 1 open reading frame 9 (Clorf9), mRNA
NM_018475	Homo sapiens TPA regulated locus (TPARL), mRNA
NM_020461	Homo sapiens gamma-tubulin complex component (GCP6), mRNA
NM_030934	Homo sapiens chromosome 1 open reading frame 25 (Clorf25), mRNA
NM_030933	Homo sapiens chromosome 1 open reading frame 14 (Clorf14), mRNA
NM 030769	Homo sapiens chromosome 1 open reading frame 13 (Clorf13), mRNA
NM 016604	Homo sapiens chromosome 5 open reading frame 7 (C5orf7), mRNA
NM_016605	Homo sapiens chromosome 5 open reading frame 6 (C5orf6), mRNA
NM 016603	Homo sapiens chromosome 5 open reading frame 5 (C5orf5), mRNA
NM_014144	Homo sapiens chromosome 11 open reading frame 21 (C11orf21), mRNA
NM_033508	Homo sapiens glucokinase (hexokinase 4, maturity onset diabetes of the young
_	2) (GCK), transcript variant 3, nuclear gene encoding mitochondrial protein, mRNA
NM 033507	Homo sapiens glucokinase (hexokinase 4, maturity onset diabetes of the young
_	2) (GCK), transcript variant 2, nuclear gene encoding mitochondrial protein, mRNA
NM 000162	Homo sapiens glucokinase (hexokinase 4, maturity onset diabetes of the young
	2) (GCK), transcript variant 1, nuclear gene encoding mitochondrial protein, mRNA
NM_025241	Homo sapiens UBX domain-containing 1 (UBXD1), mRNA
NM_002098	Homo sapiens guanylate cyclase activator 1B (retina) (GUCA1B), mRNA
NM_003137	Homo sapiens SFRS protein kinase 1 (SRPK1), mRNA
NM_003064	Homo sapiens secretory leukocyte protease inhibitor (antileukoproteinase) (SLPI), mRNA
NM 033484	Homo sapiens F-box only protein 4 (FBXO4), transcript variant 2, mRNA
	7), 44,000, 70,0000, 70,000, 70,000, 70,000, 70,000, 70,000, 70,000, 70,000, 70,000, 70,000, 70,000, 70,000, 70

NM_012176	Homo sapiens F-box only protein 4 (FBXO4), transcript variant 1, mRNA
NM_000400	Homo sapiens excision repair cross-complementing rodent repair deficiency,
	complementation group 2 (xeroderma pigmentosum D) (ERCC2), mRNA
NM_014266	Homo sapiens DNAX-activation protein 10 (DAP10), mRNA
NM_002821	Homo sapiens PTK7 protein tyrosine kinase 7 (PTK7), mRNA
NM_033502	Homo sapiens transcriptional regulating protein 132 (TReP-132), transcript
	variant 1, mRNA
NM_033501	Homo sapiens transcriptional regulating protein 132 (TReP-132), transcript
	variant 2, mRNA
NM_018415	Homo sapiens transcriptional regulating protein 132 (TReP-132), transcript
	variant 3, mRNA
NM_000994	Homo sapiens ribosomal protein L32 (RPL32), mRNA
NM_033437	Homo sapiens phosphodiesterase 5A, cGMP-specific (PDE5A), transcript variant
	3, mRNA
NM_033431	Homo sapiens phosphodiesterase 5A, cGMP-specific (PDE5A), transcript variant
	4, mRNA
NM_033430	Homo sapiens phosphodiesterase 5A, cGMP-specific (PDE5A), transcript variant
	2, mRNA
NM_001083	Homo sapiens phosphodiesterase 5A, cGMP-specific (PDE5A), transcript variant
	1, mRNA
NM_000189	Homo sapiens hexokinase 2 (HK2), mRNA
NM_033185	Homo sapiens keratin associated protein 3.3 (KAP3.3), mRNA
NM_031959	Homo sapiens keratin associated protein 3.2 (KRTAP3.2), mRNA
NM_033481	Homo sapiens F-box only protein 9 (FBXO9), transcript variant 3, mRNA
NM_033480	Homo sapiens F-box only protein 9 (FBXO9), transcript variant 2, mRNA
NM_012347	Homo sapiens F-box only protein 9 (FBXO9), transcript variant 1, mRNA
NM_033506	Homo sapiens F-box only protein 24 (FBXO24), transcript variant 1, mRNA
NM_012172	Homo sapiens F-box only protein 24 (FBXO24), transcript variant 2, mRNA
NM_012179	Homo sapiens F-box only protein 7 (FBXO7), mRNA
NM_018438	Homo sapiens F-box only protein 6 (FBXO6), mRNA
NM_012177	Homo sapiens F-box only protein 5 (FBXO5), mRNA
NM_032145	Homo sapiens F-box protein 30 (FBXO30), mRNA
NM_003813	Homo sapiens a disintegrin and metalloproteinase domain 21 (ADAM21),
ND (000014	mRNA
NM_003814	Homo sapiens a disintegrin and metalloproteinase domain 20 (ADAM20),
NB4 015600	mRNA
NM 015698	Homo sapiens T54 protein (T54), mRNA
NM_033222	Homo sapiens PC4 and SFRS1 interacting protein 2 (PSIP2), mRNA
NM_002887	Homo sapiens arginyl-tRNA synthetase (RARS), mRNA
NM 033084	Homo sapiens Fanconi anemia, complementation group D2 (FANCD2), mRNA
NM_014005 NM_018902	Homo sapiens protocadherin alpha 9 (PCDHA9), transcript variant 2, mRNA
NM 031882	Homo sapiens protocadherin alpha 11 (PCDHA11), transcript variant 1, mRNA
11111_031002	Homo sapiens protocadherin alpha subfamily C, 1 (PCDHAC1), transcript variant 2, mRNA
NM 018898	Homo sapiens protocadherin alpha subfamily C, 1 (PCDHAC1), transcript
14141_019999	variant 1, mRNA
NM 031883	
141AT_02 1 0 0 2	Homo sapiens protocadherin alpha subfamily C, 2 (PCDHAC2), transcript variant 2, mRNA
NM 018899	Homo sapiens protocadherin alpha subfamily C, 2 (PCDHAC2), transcript
144_010099	variant 1, mRNA
NM_019119	Homo sapiens protocadherin beta 9 (PCDHB9), mRNA
NM 018916	
TATAT 010310	Homo sapiens protocadherin gamma subfamily A, 3 (PCDHGA3), transcript

··	variant 1, mRNA
NM 032704	Homo sapiens tubulin alpha 6 (TUBA6), mRNA
NM_032407	Homo sapiens protocadherin gamma subfamily C, 5 (PCDHGC5), transcript variant 2, mRNA
NM_018929	Homo sapiens protocadherin gamma subfamily C, 5 (PCDHGC5), transcript variant 1, mRNA
NM_032406	Homo sapiens protocadherin gamma subfamily C, 4 (PCDHGC4), transcript variant 2, mRNA
NM_018928	Homo sapiens protocadherin gamma subfamily C, 4 (PCDHGC4), transcript variant 1, mRNA
NM_032101	Homo sapiens protocadherin gamma subfamily B, 7 (PCDHGB7), transcript variant 2, mRNA
NM_018927	Homo sapiens protocadherin gamma subfamily B, 7 (PCDHGB7), transcript variant 1, mRNA
NM_032099	Homo sapiens protocadherin gamma subfamily B, 5 (PCDHGB5), transcript variant 2, mRNA
NM_018925	Homo sapiens protocadherin gamma subfamily B, 5 (PCDHGB5), transcript variant 1, mRNA
NM_032100	Homo sapiens protocadherin gamma subfamily B, 6 (PCDHGB6), transcript variant 2, mRNA
NM_018926	Homo sapiens protocadherin gamma subfamily B, 6 (PCDHGB6), transcript variant 1, mRNA
NM_032097	Homo sapiens protocadherin gamma subfamily B, 3 (PCDHGB3), transcript variant 2, mRNA
NM_018924	Homo sapiens protocadherin gamma subfamily B, 3 (PCDHGB3), transcript variant 1, mRNA
NM_032096	Homo sapiens protocadherin gamma subfamily B, 2 (PCDHGB2), transcript variant 2, mRNA
NM_018923	Homo sapiens protocadherin gamma subfamily B, 2 (PCDHGB2), transcript variant 1, mRNA
NM_032095	Homo sapiens protocadherin gamma subfamily B, 1 (PCDHGB1), transcript variant 2, mRNA
NM_018922	Homo sapiens protocadherin gamma subfamily B, 1 (PCDHGB1), transcript variant 1, mRNA
NM_032089	Homo sapiens protocadherin gamma subfamily A, 9 (PCDHGA9), transcript variant 2, mRNA
NM_018921	Homo sapiens protocadherin gamma subfamily A, 9 (PCDHGA9), transcript variant 1, mRNA
NM_032088	Homo sapiens protocadherin gamma subfamily A, 8 (PCDHGA8), transcript variant 1, mRNA
NM_014004	Homo sapiens protocadherin gamma subfamily A, 8 (PCDHGA8), transcript variant 2, mRNA
NM_032853	Homo sapiens hypothetical protein FLJ14868 (FLJ14868), mRNA
NM_032589	Homo sapiens Down syndrome critical region gene 8 (DSCR8), mRNA
NM_032087	Homo sapiens protocadherin gamma subfamily A, 7 (PCDHGA7), transcript variant 2, mRNA
NM_018920	Homo sapiens protocadherin gamma subfamily A, 7 (PCDHGA7), transcript variant 1, mRNA
NM_032086	Homo sapiens protocadherin gamma subfamily A, 6 (PCDHGA6), transcript variant 2, mRNA
NM_018919	Homo sapiens protocadherin gamma subfamily A, 6 (PCDHGA6), transcript variant 1, mRNA

NM_032054	Homo sapiens protocadherin gamma subfamily A, 5 (PCDHGA5), transcript variant 2, mRNA
NM_018918	Homo sapiens protocadherin gamma subfamily A, 5 (PCDHGA5), transcript variant 1, mRNA
NM_032053	Homo sapiens protocadherin gamma subfamily A, 4 (PCDHGA4), transcript variant 2, mRNA
NM_018917	Homo sapiens protocadherin gamma subfamily A, 4 (PCDHGA4), transcript variant 1, mRNA
NM_032011	Homo sapiens protocadherin gamma subfamily A, 3 (PCDHGA3), transcript variant 2, mRNA
NM_032009	Homo sapiens protocadherin gamma subfamily A, 2 (PCDHGA2), transcript variant 2, mRNA
NM_018915	Homo sapiens protocadherin gamma subfamily A, 2 (PCDHGA2), transcript variant 1, mRNA
NM_031993	Homo sapiens protocadherin gamma subfamily A, 1 (PCDHGA1), transcript variant 2, mRNA
NM_032092	Homo sapiens protocadherin gamma subfamily A, 11 (PCDHGA11), transcript variant 3, mRNA
NM_018912	Homo sapiens protocadherin gamma subfamily A, 1 (PCDHGA1), transcript variant 1, mRNA
NM_032091	Homo sapiens protocadherin gamma subfamily A, 11 (PCDHGA11), transcript variant 2, mRNA
NM_018914	Homo sapiens protocadherin gamma subfamily A, 11 (PCDHGA11), transcript variant 1, mRNA
NM_032090	Homo sapiens protocadherin gamma subfamily A, 10 (PCDHGA10), transcript variant 2, mRNA
NM_018913	Homo sapiens protocadherin gamma subfamily A, 10 (PCDHGA10), transcript variant 1, mRNA
NM 019120	Homo sapiens protocadherin beta 8 (PCDHB8), mRNA
NM 018940	Homo sapiens protocadherin beta 7 (PCDHB7), mRNA
NM 018939	Homo sapiens protocadherin beta 6 (PCDHB6), mRNA
NM 015669	Homo sapiens protocadherin beta 5 (PCDHB5), mRNA
NM 018938	Homo sapiens protocadherin beta 4 (PCDHB4), mRNA
NM 018937	Homo sapiens protocadherin beta 3 (PCDHB3), mRNA
NM_018936	Homo sapiens protocadherin beta 2 (PCDHB2), mRNA
NM 013340	
	Homo sapiens protocadherin beta 1 (PCDHB1), mRNA
NM_020957	Homo sapiens protocadherin beta 16 (PCDHB16), mRNA
NM_018935	Homo sapiens protocadherin beta 15 (PCDHB15), mRNA
NM_018934	Homo sapiens protocadherin beta 14 (PCDHB14), mRNA
NM_018933	Homo sapiens protocadherin beta 13 (PCDHB13), mRNA
NM_018932	Homo sapiens protocadherin beta 12 (PCDHB12), mRNA
NM_018931	Homo sapiens protocadherin beta 11 (PCDHB11), mRNA
NM_018930	Homo sapiens protocadherin beta 10 (PCDHB10), mRNA
NM_031857	Homo sapiens protocadherin alpha 9 (PCDHA9), transcript variant 1, mRNA
NM_031856	Homo sapiens protocadherin alpha 8 (PCDHA8), transcript variant 2, mRNA
NM_018911	Homo sapiens protocadherin alpha 8 (PCDHA8), transcript variant 1, mRNA
NM_031852	Homo sapiens protocadherin alpha 7 (PCDHA7), transcript variant 2, mRNA
NM_018910	Homo sapiens protocadherin alpha 7 (PCDHA7), transcript variant 1, mRNA
NM_031501	Homo sapiens protocadherin alpha 5 (PCDHA5), transcript variant 2, mRNA
NM_018908	Homo sapiens protocadherin alpha 5 (PCDHA5), transcript variant 1, mRNA
NM_031500	Homo sapiens protocadherin alpha 4 (PCDHA4), transcript variant 2, mRNA
NM_018907	Homo sapiens protocadherin alpha 4 (PCDHA4), transcript variant 1, mRNA
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NM_031497	Homo sapiens protocadherin alpha 3 (PCDHA3), transcript variant 2, mRNA
NM_018906	Homo sapiens protocadherin alpha 3 (PCDHA3), transcript variant 1, mRNA
NM_031496	Homo sapiens protocadherin alpha 2 (PCDHA2), transcript variant 3, mRNA
NM_031495	Homo sapiens protocadherin alpha 2 (PCDHA2), transcript variant 2, mRNA
NM_018905	Homo sapiens protocadherin alpha 2 (PCDHA2), transcript variant 1, mRNA
NM_031411	Homo sapiens protocadherin alpha 1 (PCDHA1), transcript variant 3, mRNA
NM_031410	Homo sapiens protocadherin alpha 1 (PCDHA1), transcript variant 2, mRNA
NM_018900	Homo sapiens protocadherin alpha 1 (PCDHA1), transcript variant 1, mRNA
NM_031865	Homo sapiens protocadherin alpha 13 (PCDHA13), transcript variant 2, mRNA
NM_018904	Homo sapiens protocadherin alpha 13 (PCDHA13), transcript variant 1, mRNA
NM_031849	Homo sapiens protocadherin alpha 6 (PCDHA6), transcript variant 3, mRNA
NM_031864	Homo sapiens protocadherin alpha 12 (PCDHA12), transcript variant 2, mRNA
NM_031848	Homo sapiens protocadherin alpha 6 (PCDHA6), transcript variant 2, mRNA
NM_018903	Homo sapiens protocadherin alpha 12 (PCDHA12), transcript variant 1, mRNA
NM_031861	Homo sapiens protocadherin alpha 11 (PCDHA11), transcript variant 2, mRNA
NM_018909	Homo sapiens protocadherin alpha 6 (PCDHA6), transcript variant 1, mRNA
NM_031860	Homo sapiens protocadherin alpha 10 (PCDHA10), transcript variant 3, mRNA
NM_031859	Homo sapiens protocadherin alpha 10 (PCDHA10), transcript variant 2, mRNA
NM_018901	Homo sapiens protocadherin alpha 10 (PCDHA10), transcript variant 1, mRNA
NM_015429	Homo sapiens DKFZP586L2024 protein (NESHBP), mRNA
NM_031481	Homo sapiens solute carrier family 25, (mitochondrial carrier), member 18
	(SLC25A18), mRNA
NM_031442	Homo sapiens brain cell membrane protein 1 (BCMP1), mRNA
NM_030762	Homo sapiens basic helix-loop-helix domain containing, class B, 3 (BHLHB3),
	mRNA
NM_023035	Homo sapiens calcium channel, voltage-dependent, P/Q type, alpha 1A subunit
	(CACNA1A), transcript variant 2, mRNA
NM_014487	Homo sapiens nucleolar cysteine-rich protein (HSA6591), mRNA
NM_025239	Homo sapiens programmed death ligand 2 (PDL2), mRNA
NM_024859	Homo sapiens hypothetical protein FLJ21687 (FLJ21687), mRNA
NM_000575	Homo sapiens interleukin 1, alpha (IL1A), mRNA
NM_005348	Homo sapiens heat shock 90kD protein 1, alpha (HSPCA), mRNA
NM_006900	Homo sapiens interferon, alpha 13 (IFNA13), mRNA
NM_023067	Homo sapiens forkhead transcription factor FOXL2 (BPES), mRNA
NM_022552	Homo sapiens DNA (cytosine-5-)-methyltransferase 3 alpha (DNMT3A), mRNA
NM_022346	Homo sapiens chromosome condensation protein G (HCAP-G), mRNA
NM_022119	Homo sapiens protease, serine, 22 (PRSS22), mRNA
NM_022062	Homo sapiens PBX/knotted 1 homeobox 2 (PKNOX2), mRNA
NM_018665	Homo sapiens DEAD-box protein (HAGE), mRNA
NM_004614	Homo sapiens thymidine kinase 2, mitochondrial (TK2), mRNA
NM_020346	Homo sapiens solute carrier family 17 (sodium-dependent inorganic phosphate
77.4 000000	cotransporter), member 6 (SLC17A6), mRNA
NM_020309	Homo sapiens solute carrier family 17 (sodium-dependent inorganic phosphate
ND 6 000101	cotransporter), member 7 (SLC17A7), mRNA
NM_020131	Homo sapiens chromosome 1 open reading frame 6 (Clorf6), mRNA
NM_017444	Homo sapiens chromatin accessibility complex 1 (CHRAC1), mRNA
NM_016260	Homo sapiens zinc finger protein, subfamily 1A, 2 (Helios) (ZNFN1A2), mRNA
NM_015510	Homo sapiens DKFZP566O084 protein (DKFZp566O084), mRNA
NM_014433	Homo sapiens rhabdoid tumor deletion region gene 1 (RTDR1), mRNA
NM_014312	Homo sapiens cortical thymocyte receptor (X. laevis CTX) like (CTXL), mRNA
NM 004539 NM 013284	Homo sapiens asparaginyl-tRNA synthetase (NARS), mRNA Homo sapiens polymerase (DNA directed), mu (POLM), mRNA
	I Home content nelymerate (LINIA directed), mix (PCILIVI), mkina

NM_013274	Homo sapiens polymerase (DNA directed), lambda (POLL), mRNA
NM_003235	Homo sapiens thyroglobulin (TG), mRNA
NM_001963	Homo sapiens epidermal growth factor (beta-urogastrone) (EGF), mRNA
NM_007158	Homo sapiens NRAS-related gene (D1S155E), mRNA
NM_007000	Homo sapiens uroplakin 1A (UPK1A), mRNA
NM_006947	Homo sapiens signal recognition particle 72kD (SRP72), mRNA
NM_006892	Homo sapiens DNA (cytosine-5-)-methyltransferase 3 beta (DNMT3B), mRNA
NM_006760	Homo sapiens uroplakin 2 (UPK2), mRNA
NM_006691	Homo sapiens extracellular link domain-containing 1 (XLKD1), mRNA
NM_006572	Homo sapiens guanine nucleotide binding protein (G protein), alpha 13 (GNA13), mRNA
NM_006494	Homo sapiens Ets2 repressor factor (ERF), mRNA
NM_006352	Homo sapiens zinc finger protein 238 (ZNF238), mRNA
NM_006082	Homo sapiens tubulin, alpha, ubiquitous (K-ALPHA-1), mRNA
NM_005084	Homo sapiens phospholipase A2, group VII (platelet-activating factor
	acetylhydrolase, plasma) (PLA2G7), mRNA
NM_004999	Homo sapiens myosin VI (MYO6), mRNA
NM_004937	Homo sapiens cystinosis, nephropathic (CTNS), mRNA
NM_004212	Homo sapiens solute carrier family 28 (sodium-coupled nucleoside transporter), member 2 (SLC28A2), mRNA
NM_004555	Homo sapiens nuclear factor of activated T-cells, cytoplasmic, calcineurin-dependent 3 (NFATC3), mRNA
NM_004554	Homo sapiens nuclear factor of activated T-cells, cytoplasmic, calcineurin-dependent 4 (NFATC4), mRNA
NM_000695	Homo sapiens aldehyde dehydrogenase 3 family, member B2 (ALDH3B2), mRNA
NM_000373	Homo sapiens uridine monophosphate synthetase (orotate phosphoribosyl transferase and orotidine-5'-decarboxylase) (UMPS), mRNA
NM_003332	Homo sapiens TYRO protein tyrosine kinase binding protein (TYROBP), mRNA
NM 000367	Homo sapiens thiopurine S-methyltransferase (TPMT), mRNA
NM_001250	Homo sapiens tumor necrosis factor receptor superfamily, member 5 (TNFRSF5), mRNA
NM_002880	Homo sapiens v-raf-1 murine leukemia viral oncogene homolog 1 (RAF1), mRNA
NM_003978	Homo sapiens proline-serine-threonine phosphatase interacting protein 1 (PSTPIP1), mRNA
NM_003627	Homo sapiens prostate cancer overexpressed gene 1 (POV1), mRNA
NM_002557	Homo sapiens oviductal glycoprotein 1, 120kD (mucin 9, oviductin) (OVGP1), mRNA
NM_002541	Homo sapiens oxoglutarate (alpha-ketoglutarate) dehydrogenase (lipoamide) (OGDH), mRNA
NM_000406	Homo sapiens gonadotropin-releasing hormone receptor (GNRHR), mRNA
NM_001979	Homo sapiens epoxide hydrolase 2, cytoplasmic (EPHX2), mRNA
NM_001761	Homo sapiens cyclin F (CCNF), mRNA
NM_001190	Homo sapiens branched chain aminotransferase 2, mitochondrial (BCAT2), mRNA
NM_000485	Homo sapiens adenine phosphoribosyltransferase (APRT), mRNA
NM 033514	Homo sapiens pinch-2 (LOC96626), mRNA
NM 033495	Homo sapiens KIAA1309 protein (KIAA1309), mRNA
NM_022436	Homo sapiens ATP-binding cassette, sub-family G (WHITE), member 5 (sterolin
	1) (ABCG5), mRNA

NM_016333	Homo sapiens serine/arginine repetitive matrix 2 (SRRM2), mRNA
NM_012412	Homo sapiens histone H2A.F/Z variant (H2AV), mRNA
NM_001897	Homo sapiens chondroitin sulfate proteoglycan 4 (melanoma-associated)
	(CSPG4), mRNA
NM_031420	Homo sapiens mitochondrial ribosomal protein L9 (MRPL9), mRNA
NM_020393	Homo sapiens hypothetical protein SBBI67 (LOC57115), mRNA
NM_015956	Homo sapiens mitochondrial ribosomal protein L4 (MRPL4), mRNA
NM_004537	Homo sapiens nucleosome assembly protein 1-like 1 (NAP1L1), mRNA
NM_033504	Homo sapiens CAC-1 (CAC-1), mRNA
NM_033503	Homo sapiens Bcl-2 modifying factor (BMF), mRNA
NM_022059	Homo sapiens chemokine (C-X-C motif) ligand 16 (CXCL16), mRNA
NM_022048	Homo sapiens casein kinase 1, gamma 1 (CSNK1G1), mRNA
NM_019009	Homo sapiens Toll-interacting protein (TOLLIP), mRNA
NM_018058	Homo sapiens cartilage acidic protein 1 (CRTAC1), mRNA
NM_017443	Homo sapiens polymerase (DNA directed), epsilon 3 (p17 subunit) (POLE3), mRNA
NM_007359	Homo sapiens MLN51 protein (MLN51), mRNA
NM_030956	Homo sapiens toll-like receptor 10 (TLR10), mRNA
NM_020653	Homo sapiens zinc finger protein 287 (ZNF287), mRNA
NM_020652	Homo sapiens zinc finger protein 286 (ZNF286), mRNA
NM_020365	Homo sapiens eukaryotic translation initiation factor 2B, subunit 3 (gamma, 58kD) (EIF2B3), mRNA
NM_013432	Homo sapiens nuclear factor of kappa light polypeptide gene enhancer in B-cells inhibitor-like 2 (NFKBIL2), mRNA
NM_003740	Homo sapiens potassium channel, subfamily K, member 5 (TASK-2) (KCNK5), mRNA
NM_033311	Homo sapiens potassium inwardly-rectifying channel, subfamily K, member 4 (KCNK4), transcript variant 3, mRNA
NM_033310	Homo sapiens potassium inwardly-rectifying channel, subfamily K, member 4 (KCNK4), transcript variant 2, mRNA
NM_016611	Homo sapiens potassium inwardly-rectifying channel, subfamily K, member 4 (KCNK4), transcript variant 1, mRNA
NM_033360	Homo sapiens v-Ki-ras2 Kirsten rat sarcoma 2 viral oncogene homolog (KRAS2), transcript variant a, mRNA
NM_004985	Homo sapiens v-Ki-ras2 Kirsten rat sarcoma 2 viral oncogene homolog (KRAS2), transcript variant b, mRNA
NM_022442	Homo sapiens ubiquitin-conjugating enzyme E2 variant 1 (UBE2V1), transcript variant 3, mRNA
NM_021988	Homo sapiens ubiquitin-conjugating enzyme E2 variant 1 (UBE2V1), transcript variant 1, mRNA
NM_003349	Homo sapiens ubiquitin-conjugating enzyme E2 variant 1 (UBE2V1), transcript variant 2, mRNA
NM_003546	Homo sapiens H4 histone family, member K (H4FK), mRNA
NM_003541	Homo sapiens H4 histone family, member D (H4FD), mRNA
NM_003536	Homo sapiens H3 histone family, member K (H3FK), mRNA
NM_003535	Homo sapiens H3 histone family, member J (H3FJ), mRNA
NM_003533	Homo sapiens H3 histone family, member F (H3FF), mRNA
NM_003521	Homo sapiens H2B histone family, member E (H2BFE), mRNA
NM_003520	Homo sapiens H2B histone family, member D (H2BFD), mRNA
NM_003519	Homo sapiens H2B histone family, member C (H2BFC), mRNA
NM 003514	Homo sapiens H2A histone family, member N (H2AFN), mRNA
NM_003511	Homo sapiens H2A histone family, member I (H2AFI), mRNA

NM 005322	Homo sapiens H1 histone family, member 5 (H1F5), mRNA
NM 021066	Homo sapiens H2A histone family, member E (H2AFE), mRNA
NM 003510	Homo sapiens H2A histone family, member D (H2AFD), mRNA
NM 003509	Homo sapiens H2A histone family, member C (H2AFC), mRNA
NM 033358	Homo sapiens caspase 8, apoptosis-related cysteine protease (CASP8), transcript
	variant E, mRNA
NM_033357	Homo sapiens caspase 8, apoptosis-related cysteine protease (CASP8), transcript variant D, mRNA
NM_033356	Homo sapiens caspase 8, apoptosis-related cysteine protease (CASP8), transcript variant C, mRNA
NM_033355	Homo sapiens caspase 8, apoptosis-related cysteine protease (CASP8), transcript variant B, mRNA
NM_001228	Homo sapiens caspase 8, apoptosis-related cysteine protease (CASP8), transcript variant A, mRNA
NM_033340	Homo sapiens caspase 7, apoptosis-related cysteine protease (CASP7), transcript variant beta, mRNA
NM_033339	Homo sapiens caspase 7, apoptosis-related cysteine protease (CASP7), transcript variant gamma, mRNA
NM_033338	Homo sapiens caspase 7, apoptosis-related cysteine protease (CASP7), transcript variant delta, mRNA
NM_001227	Homo sapiens caspase 7, apoptosis-related cysteine protease (CASP7), transcript variant alpha, mRNA
NM_001005	Homo sapiens ribosomal protein S3 (RPS3), mRNA
NM 006013	Homo sapiens ribosomal protein L10 (RPL10), mRNA
NM 013368	Homo sapiens RPA-binding trans-activator (RBT1), mRNA
NM 002286	Homo sapiens lymphocyte-activation gene 3 (LAG3), mRNA
NM 005546	Homo sapiens IL2-inducible T-cell kinase (ITK), mRNA
NM 005538	Homo sapiens inhibin, beta C (INHBC), mRNA
NM_033257	Homo sapiens DiGeorge syndrome critical region gene 6 like (DGCR6L), mRNA
NM 001917	Homo sapiens D-amino-acid oxidase (DAO), mRNA
NM_001629	Homo sapiens arachidonate 5-lipoxygenase-activating protein (ALOX5AP), mRNA
NM 000024	Homo sapiens adrenergic, beta-2-, receptor, surface (ADRB2), mRNA
NM_000683	Homo sapiens adrenergic, alpha-2C-, receptor (ADRA2C), mRNA
NM 000682	Homo sapiens adrenergic, alpha-2B-, receptor (ADRA2B), mRNA
NM 000681	Homo sapiens adrenergic, alpha-2A-, receptor (ADRA2A), mRNA
NM 006179	Homo sapiens neurotrophin 5 (neurotrophin 4/5) (NTF5), mRNA
NM 033277	Homo sapiens lacritin (LACRT), mRNA
NM 022128	Homo sapiens ribokinase (RBSK), mRNA
NM_004823	Homo sapiens potassium channel, subfamily K, member 6 (TWIK-2) (KCNK6), mRNA
NM_002246	Homo sapiens potassium channel, subfamily K, member 3 (TASK-1) (KCNK3), mRNA
NM_032405	Homo sapiens transmembrane protease, serine 3 (TMPRSS3), transcript variant D, mRNA
NM_032404	Homo sapiens transmembrane protease, serine 3 (TMPRSS3), transcript variant C, mRNA
NM_032401	Homo sapiens transmembrane protease, serine 3 (TMPRSS3), transcript variant B, mRNA
NM_024022	Homo sapiens transmembrane protease, serine 3 (TMPRSS3), transcript variant A, mRNA

NM_016234	Homo sapiens fatty-acid-Coenzyme A ligase, long-chain 5 (FACL5), mRNA
NM 006883	Homo sapiens short stature homeobox (SHOX), transcript variant SHOXb,
1444_000003	mRNA
NM_000451	Homo sapiens short stature homeobox (SHOX), transcript variant SHOXa,
	mRNA
NM_006476	Homo sapiens ATP synthase, H+ transporting, mitochondrial F0 complex,
	subunit g (ATP5L), mRNA
NM_006356	Homo sapiens ATP synthase, H+ transporting, mitochondrial F0 complex,
	subunit d (ATP5H), mRNA
NM_024683	Homo sapiens hypothetical protein FLJ22729 (FLJ22729), mRNA
NM_033468	Homo sapiens zinc finger protein 257 (ZNF257), mRNA
NM_033453	Homo sapiens inosine triphosphatase (nucleoside triphosphate pyrophosphatase)
	(ITPA), mRNA
NM_032144	Homo sapiens RAB6C, member RAS oncogene family (RAB6C), mRNA
NM_031296	Homo sapiens RAB33B, member RAS oncogene family (RAB33B), mRNA
NM_022570	Homo sapiens C-type (calcium dependent, carbohydrate-recognition domain)
	lectin, superfamily member 12 (CLECSF12), mRNA
NM_022825	Homo sapiens porcupine (MG61), mRNA
NM_022449	Homo sapiens RAB17, member RAS oncogene family (RAB17), mRNA
NM_016322	Homo sapiens RAB14, member RAS oncogene family (RAB14), mRNA
NM_006331	Homo sapiens C2f protein (C2F), mRNA
NM_007066	Homo sapiens protein kinase (cAMP-dependent, catalytic) inhibitor gamma
	(PKIG), mRNA
NM_002732	Homo sapiens protein kinase, cAMP-dependent, catalytic, gamma (PRKACG), mRNA
NM_005055	Homo sapiens receptor-associated protein of the synapse, 43kD (RAPSN),
11111_005055	transcript variant 1, mRNA
NM 032645	Homo sapiens receptor-associated protein of the synapse, 43kD (RAPSN),
11212_002010	transcript variant 2, mRNA
NM 033305	Homo sapiens chorea acanthocytosis (CHAC), transcript variant A, mRNA
NM 015186	Homo sapiens chorea acanthocytosis (CHAC), transcript variant B, mRNA
NM 004624	Homo sapiens vasoactive intestinal peptide receptor 1 (VIPR1), mRNA
NM 030967	Homo sapiens keratin associated protein 1.1 (KRTAP1.1), mRNA
NM 015696	Homo sapiens weakly similar to glutathione peroxidase 2 (CL683), mRNA
NM 031885	Homo sapiens Bardet-Biedl syndrome 2 (BBS2), mRNA
NM 030966	Homo sapiens keratin associated protein 1.3 (KRTAP1.3), mRNA
NM_007083	Homo sapiens nudix (nucleoside diphosphate linked moiety X)-type motif 6
_	(NUDT6), mRNA
NM 013317	Homo sapiens lung type-I cell membrane-associated glycoprotein (T1A-2),
·	transcript variant 1, mRNA
NM_006474	Homo sapiens lung type-I cell membrane-associated glycoprotein (T1A-2),
	transcript variant 2, mRNA
NM_006275	Homo sapiens splicing factor, arginine/serine-rich 6 (SFRS6), mRNA
NM_016041	Homo sapiens CGI-101 protein (F-LAN-1), mRNA
NM_001954	Homo sapiens discoidin domain receptor family, member 1 (DDR1), transcript
	variant 2, mRNA
NM_013994	Homo sapiens discoidin domain receptor family, member 1 (DDR1), transcript
	variant 3, mRNA
NM_013993	Homo sapiens discoidin domain receptor family, member 1 (DDR1), transcript
	variant 1, mRNA
NM_022117	Homo sapiens cutaneous T-cell lymphoma-associated tumor antigen se20-4;
	differentially expressed nucleolar TGF-beta1 target protein (DENTT) (SE20-4),

	mRNA
NM 003048	Homo sapiens solute carrier family 9 (sodium/hydrogen exchanger), isoform 2
MM_003048	(SLC9A2), mRNA
NM 001971	Homo sapiens elastase 1, pancreatic (ELA1), mRNA
NM 033412	Homo sapiens hypothetical protein similar to CG7943 (MGC14836), mRNA
NM 033420	Homo sapiens hypothetical protein MGC4022 (R32184_3), mRNA
	Homo sapiens hypothetical protein MBC3205 (MBC3205), mRNA
NM_033408	Homo sapiens dual adaptor of phosphotyrosine and 3-phosphoinositides
NM_014395	(DAPP1), mRNA
NM 003918	Homo sapiens glycogenin 2 (GYG2), mRNA
NM 001502	Homo sapiens glycoprotein 2 (zymogen granule membrane) (GP2), mRNA
NM 006362	Homo sapiens nuclear RNA export factor 1 (NXF1), mRNA
NM 033155	Homo sapiens nuclear RNA export factor 5 (NXF5), transcript variant 5, mRNA
NM 033154	Homo sapiens nuclear RNA export factor 5 (NXF5), transcript variant 4, mRNA
NM 033153	Homo sapiens nuclear RNA export factor 5 (NXF5), transcript variant 3, mRNA
NM 033152	Homo sapiens nuclear RNA export factor 5 (NXF5), transcript variant 2, mRNA
NM 032946	Homo sapiens nuclear RNA export factor 5 (NXF5), transcript variant 1, mRNA
NM 022052	Homo sapiens nuclear RNA export factor 3 (NXF3), mRNA
NM 021808	Homo sapiens UDP-N-acetyl-alpha-D-galactosamine:polypeptide N-
1122	acetylgalactosaminyltransferase 9 (GalNAc-T9) (GALNT9), mRNA
NM 017840	Homo sapiens mitochondrial ribosomal protein L16 (MRPL16), mRNA
NM 017417	Homo sapiens UDP-N-acetyl-alpha-D-galactosamine:polypeptide N-
14112_017.117	acetylgalactosaminyltransferase 8 (GalNAc-T8) (GALNT8), mRNA
NM 004261	Homo sapiens 15 kDa selenoprotein (SEP15), mRNA
NM 021998	Homo sapiens zinc finger protein 6 (CMPX1) (ZNF6), mRNA
NM_004570	Homo sapiens phosphoinositide-3-kinase, class 2, gamma polypeptide
1111_00 1070	(PIK3C2G), mRNA
NM 002646	Homo sapiens phosphoinositide-3-kinase, class 2, beta polypeptide (PIK3C2B),
	mRNA
NM 004598	Homo sapiens sparc/osteonectin, cwcv and kazal-like domains proteoglycan
	(testican) (SPOCK), mRNA
NM 033135	Homo sapiens spinal cord-derived growth factor-B (SCDGF-B), transcript
	variant 2, mRNA
NM 025208	Homo sapiens spinal cord-derived growth factor-B (SCDGF-B), transcript
	variant 1, mRNA
NM_033346	Homo sapiens bone morphogenetic protein receptor, type II (serine/threonine
_	kinase) (BMPR2), transcript variant 2, mRNA
NM 001204	Homo sapiens bone morphogenetic protein receptor, type II (serine/threonine
_	kinase) (BMPR2), transcript variant 1, mRNA
NM 003933	Homo sapiens BAI1-associated protein 3 (BAIAP3), mRNA
NM_005467	Homo sapiens N-acetylated alpha-linked acidic dipeptidase 2 (NAALAD2),
-	mRNA
NM_005944	Homo sapiens antigen identified by monoclonal antibody MRC OX-2 (MOX2),
-	mRNA
NM_002245	Homo sapiens potassium channel, subfamily K, member 1 (TWIK-1) (KCNK1),
ND 6 005045	mRNA
NM_005247	Homo sapiens fibroblast growth factor 3 (murine mammary tumor virus
37.6.0000	integration site (v-int-2) oncogene homolog) (FGF3), mRNA
NM_002006	Homo sapiens fibroblast growth factor 2 (basic) (FGF2), mRNA
NM_000647	Homo sapiens chemokine (C-C motif) receptor 2 (CCR2), transcript variant A, mRNA
NM 032047	Homo sapiens UDP-GlcNAc:betaGal beta-1,3-N-acetylglucosaminyltransferase

	5 (B3GNT5), mRNA
NM_014256	Homo sapiens UDP-GlcNAc:betaGal beta-1,3-N-acetylglucosaminyltransferase
	3 (B3GNT3), mRNA
NM_015904	Homo sapiens translation initiation factor IF2 (IF2), mRNA
NM_005326	Homo sapiens hydroxyacyl glutathione hydrolase (HAGH), mRNA
NM_013445	Homo sapiens glutamate decarboxylase 1 (brain, 67kD) (GAD1), transcript
NR 6 022172	variant GAD25, mRNA
NM_033173	Homo sapiens UDP-Gal:betaGlcNAc beta 1,3-galactosyltransferase, polypeptide 5 (B3GALT5), transcript variant 5, mRNA
NM_033172	Homo sapiens UDP-Gal:betaGlcNAc beta 1,3-galactosyltransferase, polypeptide
_	5 (B3GALT5), transcript variant 4, mRNA
NM_033171	Homo sapiens UDP-Gal:betaGlcNAc beta 1,3-galactosyltransferase, polypeptide
	5 (B3GALT5), transcript variant 3, mRNA
NM_033170	Homo sapiens UDP-Gal:betaGlcNAc beta 1,3-galactosyltransferase, polypeptide
	5 (B3GALT5), transcript variant 2, mRNA
NM_033169	Homo sapiens UDP-Gal:betaGlcNAc beta 1,3-galactosyltransferase, polypeptide
	3 (B3GALT3), transcript variant 4, mRNA
NM_033168	Homo sapiens UDP-Gal:betaGlcNAc beta 1,3-galactosyltransferase, polypeptide
	3 (B3GALT3), transcript variant 3, mRNA
NM_033167	Homo sapiens UDP-Gal:betaGlcNAc beta 1,3-galactosyltransferase, polypeptide
	3 (B3GALT3), transcript variant 2, mRNA
NM_003781	Homo sapiens UDP-Gal:betaGlcNAc beta 1,3-galactosyltransferase, polypeptide
	3 (B3GALT3), transcript variant 1, mRNA
NM_003782	Homo sapiens UDP-Gal:betaGlcNAc beta 1,3-galactosyltransferase, polypeptide
	4 (B3GALT4), mRNA
NM_003783	Homo sapiens UDP-Gal:betaGlcNAc beta 1,3-galactosyltransferase, polypeptide
	2 (B3GALT2), mRNA
NM_004631	Homo sapiens low density lipoprotein receptor-related protein 8, apolipoprotein
	e receptor (LRP8), transcript variant 1, mRNA
NM_033300	Homo sapiens low density lipoprotein receptor-related protein 8, apolipoprotein
)7 f 015500	e receptor (LRP8), transcript variant 2, mRNA
NM_017522	Homo sapiens low density lipoprotein receptor-related protein 8, apolipoprotein
) D f 000000	e receptor (LRP8), transcript variant 3, mRNA
NM_033323	Homo sapiens sodium bicarbonate transporter 4 (NBC4), transcript variant b, mRNA
NM_033337	Homo sapiens caveolin 3 (CAV3), transcript variant 1, mRNA
NM_001234	Homo sapiens caveolin 3 (CAV3), transcript variant 2, mRNA
NM 001233	Homo sapiens caveolin 2 (CAV2), mRNA
NM 001753	Homo sapiens caveolin 1, caveolae protein, 22kD (CAV1), mRNA
NM 033291	Homo sapiens midline 1 (Opitz/BBB syndrome) (MID1), transcript variant 2,
	mRNA
NM_033290	Homo sapiens midline 1 (Opitz/BBB syndrome) (MID1), transcript variant 3,
	mRNA
NM 033274	Homo sapiens a disintegrin and metalloproteinase domain 19 (meltrin beta)
	(ADAM19), transcript variant 2, mRNA
NM 023038	Homo sapiens a disintegrin and metalloproteinase domain 19 (meltrin beta)
_	(ADAM19), transcript variant 1, mRNA
NM_033308	Homo sapiens ATP-binding cassette, sub-family A (ABC1), member 7
_	(ABCA7), transcript variant 2, mRNA
NM_019112	Homo sapiens ATP-binding cassette, sub-family A (ABC1), member 7
_	(ABCA7), transcript variant 1, mRNA
NM 002609	Homo sapiens platelet-derived growth factor receptor, beta polypeptide

	(MDCERD) DMA
) Tr (00/00/	(PDGFRB), mRNA
NM_006206	Homo sapiens platelet-derived growth factor receptor, alpha polypeptide
ND4 022016	(PDGFRA), mRNA Homo sapiens platelet-derived growth factor beta polypeptide (simian sarcoma
NM_033016	viral (v-sis) oncogene homolog) (PDGFB), transcript variant 2, mRNA
ND4 000679	
NM_000678	Homo sapiens adrenergic, alpha-1D-, receptor (ADRA1D), mRNA
NM_000679	Homo sapiens adrenergic, alpha-1B-, receptor (ADRA1B), mRNA
NM_002675	Homo sapiens promyelocytic leukemia (PML), transcript variant 6, mRNA
NM_033250	Homo sapiens promyelocytic leukemia (PML), transcript variant 11, mRNA
NM_033249	Homo sapiens promyelocytic leukemia (PML), transcript variant 10, mRNA
NM_033247	Homo sapiens promyelocytic leukemia (PML), transcript variant 8, mRNA
NM_033246	Homo sapiens promyelocytic leukemia (PML), transcript variant 7, mRNA
NM_033245	Homo sapiens promyelocytic leukemia (PML), transcript variant 12, mRNA
NM_033244	Homo sapiens promyelocytic leukemia (PML), transcript variant 5, mRNA
NM_033242	Homo sapiens promyelocytic leukemia (PML), transcript variant 3, mRNA
NM_033240	Homo sapiens promyelocytic leukemia (PML), transcript variant 2, mRNA
NM_033239	Homo sapiens promyelocytic leukemia (PML), transcript variant 9, mRNA
NM_033238	Homo sapiens promyelocytic leukemia (PML), transcript variant 1, mRNA
NM_033304	Homo sapiens adrenergic, alpha-1A-, receptor (ADRA1A), transcript variant 4, mRNA
NM_033303	Homo sapiens adrenergic, alpha-1A-, receptor (ADRA1A), transcript variant 2, mRNA
NM_033302	Homo sapiens adrenergic, alpha-1A-, receptor (ADRA1A), transcript variant 3, mRNA
NM 033279	Homo sapiens ring finger protein 22 (RNF22), transcript variant gamma, mRNA
NM 033278	Homo sapiens ring finger protein 22 (RNF22), transcript variant beta, mRNA
NM 000737	Homo sapiens chorionic gonadotropin, beta polypeptide (CGB), mRNA
NM_033295	Homo sapiens caspase 1, apoptosis-related cysteine protease (interleukin 1, beta,
_	convertase) (CASP1), transcript variant epsilon, mRNA,
NM 033294	Homo sapiens caspase 1, apoptosis-related cysteine protease (interleukin 1, beta,
_	convertase) (CASP1), transcript variant delta, mRNA
NM_033293	Homo sapiens caspase 1, apoptosis-related cysteine protease (interleukin 1, beta,
_	convertase) (CASP1), transcript variant gamma, mRNA
NM_033292	Homo sapiens caspase 1, apoptosis-related cysteine protease (interleukin 1, beta,
_	convertase) (CASP1), transcript variant alpha, mRNA
NM_001223	Homo sapiens caspase 1, apoptosis-related cysteine protease (interleukin 1, beta,
_	convertase) (CASP1), transcript variant beta, mRNA
NM_006771	Homo sapiens keratin, hair, acidic, 8 (KRTHA8), mRNA
NM 002280	Homo sapiens keratin, hair, acidic, 5 (KRTHA5), mRNA
NM 000526	Homo sapiens keratin 14 (epidermolysis bullosa simplex, Dowling-Meara,
_	Koebner) (KRT14), mRNA
NM 033301	Homo sapiens ribosomal protein L8 (RPL8), transcript variant 2, mRNA
NM 000973	Homo sapiens ribosomal protein L8 (RPL8), franscript variant 1, mRNA
NM_000661	Homo sapiens ribosomal protein L9 (RPL9), mRNA
NM 007104	Homo sapiens ribosomal protein L10a (RPL10A), mRNA
NM 033255	Homo sapiens epithelial stromal interaction 1 (breast) (EPSTI1), mRNA
NM_021196	Homo sapiens sodium bicarbonate transporter 4 (NBC4), transcript variant a, mRNA
NM 032241	Homo sapiens ribosomal protein L10 (RPL10), mRNA
NM 030955	Homo sapiens a disintegrin-like and metalloprotease (reprolysin type) with
	thrombospondin type 1 motif, 12 (ADAMTS12), mRNA
NM_030765	Homo sapiens UDP-GlcNAc:betaGal beta-1,3-N-acetylglucosaminyltransferase

r	L COOCHEAN BY
L	4 (B3GNT4), mRNA
NM_014670	Homo sapiens basic leucine-zipper protein BZAP45 (BZAP45), mRNA
NM_013379	Homo sapiens dipeptidylpeptidase 7 (DPP7), mRNA
NM_006458	Homo sapiens ring finger protein 22 (RNF22), transcript variant alpha, mRNA
NM_006057	Homo sapiens UDP-Gal:betaGlcNAc beta 1,3-galactosyltransferase, polypeptide 5 (B3GALT5), transcript variant 1, mRNA
NM_000648	Homo sapiens chemokine (C-C motif) receptor 2 (CCR2), transcript variant B, mRNA
NM_000381	Homo sapiens midline 1 (Opitz/BBB syndrome) (MID1), transcript variant 1, mRNA
NM_002645	Homo sapiens phosphoinositide-3-kinase, class 2, alpha polypeptide (PIK3C2A), mRNA
NM_002608	Homo sapiens platelet-derived growth factor beta polypeptide (simian sarcoma viral (v-sis) oncogene homolog) (PDGFB), transcript variant 1, mRNA
NM 001134	Homo sapiens alpha-fetoprotein (AFP), mRNA
NM_000680	Homo sapiens adrenergic, alpha-1A-, receptor (ADRA1A), transcript variant 1, mRNA
NM 023929	Homo sapiens zinc finger protein RINZF (RINZF), mRNA
NM 020353	Homo sapiens phospholipid scramblase 4 (PLSCR4), mRNA
NM 020359	Homo sapiens phospholipid scramblase 2 (PLSCR2), mRNA
NM 018494	Homo sapiens leucine-rich and death domain containing (LRDD), mRNA
NM 004998	Homo sapiens myosin IE (MYO1E), mRNA
NM 033226	Homo sapiens ATP-binding cassette, sub-family C (CFTR/MRP), member 12
	(ABCC12), mRNA
NM_032105	Homo sapiens protein phosphatase 1, regulatory (inhibitor) subunit 12B (PPP1R12B), transcript variant 2, mRNA
NM_032104	Homo sapiens protein phosphatase 1, regulatory (inhibitor) subunit 12B (PPP1R12B), transcript variant 4, mRNA
NM_032103	Homo sapiens protein phosphatase 1, regulatory (inhibitor) subunit 12B (PPP1R12B), transcript variant 3, mRNA
NM_002481	Homo sapiens protein phosphatase 1, regulatory (inhibitor) subunit 12B (PPP1R12B), transcript variant 1, mRNA
NM 004689	Homo sapiens metastasis associated 1 (MTA1), mRNA
NM 006005	Homo sapiens Wolfram syndrome 1 (wolframin) (WFS1), mRNA
NM_015722	Homo sapiens calcyon; D1 dopamine receptor-interacting protein (CALCYON), mRNA
NM 004184	Homo sapiens tryptophanyl-tRNA synthetase (WARS), mRNA
NM_014228	Homo sapiens solute carrier family 6 (neurotransmitter transporter, L-proline), member 7 (SLC6A7), mRNA
NM 005823	Homo sapiens mesothelin (MSLN), transcript variant 1, mRNA
NM 013404	Homo sapiens mesothelin (MSLN), transcript variant 2, mRNA
NM 012341	Homo sapiens G protein-binding protein CRFG (CRFG), mRNA
NM_002480	Homo sapiens protein phosphatase 1, regulatory (inhibitor) subunit 12A (PPP1R12A), mRNA
NM 003868	Homo sapiens fibroblast growth factor 16 (FGF16), mRNA
NM 018979	Homo sapiens protein kinase, lysine deficient 1 (PRKWNK1), mRNA
NM_022127	Homo sapiens solute carrier family 28 (sodium-coupled nucleoside transporter), member 3 (SLC28A3), mRNA
NM_005517	Homo sapiens high-mobility group (nonhistone chromosomal) protein 17 (HMG17), mRNA
NM 022465	Homo sapiens zinc finger protein, subfamily 1A, 4 (Eos) (ZNFN1A4), mRNA
NM 005768	Homo sapiens putative protein similar to nessy (Drosophila) (C3F), mRNA
1111 003 / 08	1 Mondo Suprano pulatiro protein similar to nossy (Dissoprina) (OSI), Intern

NM 033199	Homo sapiens stresscopin-related peptide (SRP), mRNA
NM_032243	Homo sapiens thioredoxin domain-containing 2 (spermatozoa) (TXNDC2),
11111_032243	mRNA
NM 031433	Homo sapiens membrane-type frizzled-related protein (MFRP), mRNA
NM 022466	Homo sapiens zinc finger protein, subfamily 1A, 5 (Pegasus) (PEGASUS),
NWI_022400	mRNA
NM_004320	Homo sapiens ATPase, Ca++ transporting, cardiac muscle, fast twitch 1
	(ATP2A1), mRNA
NM_021047	Homo sapiens zinc finger protein 253 (ZNF253), mRNA
NM_020152	Homo sapiens chromosome 21 open reading frame 7 (C21orf7), mRNA
NM_017447	Homo sapiens chromosome 21 open reading frame 91 (C21orf91), mRNA
NM_016154	Homo sapiens RAB4B, member RAS oncogene family (RAB4B), mRNA
NM_016308	Homo sapiens UMP-CMP kinase (UMP-CMPK), mRNA
NM_016066	Homo sapiens glutaredoxin 2 (GLRX2), mRNA
NM_016255	Homo sapiens family with sequence similarity 8, member A1 (FAM8A1), mRNA
ND4 014701	Homo sapiens likely ortholog of mouse coiled coil forming protein 1
NM_014781	(KIAA0203), mRNA
NM 014468	Homo sapiens VENT-like homeobox 2 (VENTX2), mRNA
NM 013383	Homo sapiens transcription factor-like 4 (TCFL4), mRNA
NM 012481	Homo sapiens zinc finger protein, subfamily 1A, 3 (Aiolos) (ZNFN1A3), mRNA
NM 012230	Homo sapiens POM (POM121 rat homolog) and ZP3 fusion (POMZP3), mRNA
NM 012199	Homo sapiens eukaryotic translation initiation factor 2C, 1 (EIF2C1), mRNA
NM 005849	Homo sapiens immunoglobulin superfamily, member 6 (IGSF6), mRNA
NM_005414	Homo sapiens SKI-like (SKIL), mRNA
NM 004245	Homo sapiens transglutaminase 5 (TGM5), mRNA
NM 020831	Homo sapiens megakaryoblastic leukemia (translocation) 1 (MKL1), mRNA
NM 015870	Homo sapiens endogenous retrovirus H D1 leader region/integrase-derived
11112_015070	ORF1, ORF2, and putative envelope protein (HSU88895), mRNA
NM 033330	Homo sapiens scavenger receptor cysteine-rich type 1 protein M160 precursor
- ···· -	(M160), mRNA
NM 033326	Homo sapiens Sox-6 (HSSOX6), mRNA
NM 017829	Homo sapiens cat eye syndrome chromosome region, candidate 5 (CECR5),
_	mRNA
NM_033256	Homo sapiens protein phosphatase 1, regulatory (inhibitor) subunit 14A
	(PPP1R14A), mRNA
NM_033213	Homo sapiens hypothetical protein MGC12466 (MGC12466), mRNA
NM_033070	Homo sapiens cat eye syndrome chromosome region, candidate 5 (CECR5),
	mRNA
NM_032752	Homo sapiens hypothetical protein MGC15548 (MGC15548), mRNA
NM_032686	Homo sapiens hypothetical protein MGC13008 (MGC13008), mRNA
NM_032371	Homo sapiens hypothetical protein MGC15416 (MGC15416), mRNA
NM_032366	Homo sapiens hypothetical protein MGC13114 (MGC13114), mRNA
NM_032353	Homo sapiens hypothetical protein MGC10540 (MGC10540), mRNA
NM_032304	Homo sapiens hypothetical protein MGC2605 (MGC2605), mRNA
NM_032259	Homo sapiens hypothetical protein DKFZp434F054 (DKFZp434F054), mRNA
NM_032240	Homo sapiens hypothetical protein FLJ23519 (FLJ23519), mRNA
NM_032153	Homo sapiens zinc family member 4 protein HZIC4 (ZIC4), mRNA
NM_015064	Homo sapiens ELKS protein (ELKS), mRNA
NM_031294	Homo sapiens hypothetical protein DKFZp586M1120 (DKFZP586M1120), mRNA
NM_025213	Homo sapiens spectrin, beta, non-erythrocytic 4 (SPTBN4), mRNA

NM 025267 Homo sapiens hypothetical protein FIJ23022 (FIJ23022), mRNA NM 024974 Homo sapiens hypothetical protein FIJ23022 (FIJ23022), mRNA NM 024934 Homo sapiens hypothetical protein FIJ211800 (FIJ211800), mRNA NM 024935 Homo sapiens hypothetical protein FIJ21172 (FIJ21172), mRNA NM 024806 Homo sapiens hypothetical protein FIJ212659 (FIJ22659), mRNA NM 024807 Homo sapiens hypothetical protein FIJ21172 (FIJ21172), mRNA NM 024808 Homo sapiens hypothetical protein FIJ21172 (FIJ21172), mRNA NM 024042 Homo sapiens hypothetical protein MGC3048 (MGC3048), mRNA NM 024042 Homo sapiens hypothetical protein MGC2601 (MGC2601), mRNA NM 024042 Homo sapiens hypothetical protein MGC2601 (MGC2601), mRNA NM 021939 Homo sapiens hypothetical protein FIJ21041 similar to FK506 binding proteins (FIJ22041), mRNA NM 018722 Homo sapiens BWRT protein (HSA404617), mRNA NM 018722 Homo sapiens BWRT protein (HSA404617), mRNA NM 018724 Homo sapiens BWRT protein (HSA404617), mRNA NM 018629 Homo sapiens super notein SER73 (LOC57116), mRNA NM 018629 Homo sapiens hypothetical protein FR02533 (PR02533), mRNA NM 018636 Homo sapiens hypothetical protein FR02533 (PR02533), mRNA NM 018019 Homo sapiens hypothetical protein FR110193 (FR03043), mRNA NM 018050 Homo sapiens hypothetical protein FL110298 (FL110298), mRNA NM 016332 Homo sapiens hypothetical protein FL10298 (FL110298), mRNA NM 016330 Homo sapiens hypothetical protein FL10298 (FL110298), mRNA NM 016330 Homo sapiens selenoprotein X, 1 (SEPX1), mRNA NM 016360 Homo sapiens SIAA0663 protein (KIAA0663), mRNA NM 016360 Homo sapiens SIAA0663 protein (KIAA0665), mRNA NM 016480 Homo sapiens SIAA0663 protein (KIAA0665), mRNA NM 014700 Homo sapiens SIAA0663 protein (KIAA0665), mRNA NM 014680 Homo sapiens SIAA0663 protein (KIAA0665), mRNA NM 014680 Homo sapiens SIAA0665 protein (KIAA0665), mRNA NM 014680 Homo sapiens SIAA0665 protein (KIAA0665), mRNA NM 016603 Homo sapiens SIAA0100 gene product (KIAA0100), mRNA NM 016604 Homo sapiens SIAA0665 protein (KIAA0665), mRNA NM 006653 Homo sapiens SIAA0665 prote		the state of the s
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NM 019013		Homo sapiens BWRT protein (HSA404617), mRNA
NM 018629		Homo sapiens zinc finger protein SBZF3 (LOC57116), mRNA
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NM_005675 Homo sapiens DiGeorge syndrome critical region gene 6 (DGCR6), mRNA NM_016083 Homo sapiens cannabinoid receptor 1 (brain) (CNR1), transcript variant 2, mRNA NM_004053 Homo sapiens bystin-like (BYSL), mRNA NG_000016 Homo sapi ns genomic protocadherin alpha cluster (PCDHA@) on chromosome 5	NM_033023	Homo sapiens platelet-derived growth factor alpha polypeptide (PDGFA),
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NM_004053 Homo sapiens bystin-like (BYSL), mRNA NG_000016 Homo sapi ns genomic protocadherin alpha cluster (PCDHA@) on chromosome 5		Homo sapiens cannabinoid receptor 1 (brain) (CNR1), transcript variant 2,
NG_000016 Homo sapi ns genomic protocadherin alpha cluster (PCDHA@) on chromosome 5	NM 004053	Homo sapiens bystin-like (BYSL), mRNA
		Homo sapi ns genomic protocadherin alpha cluster (PCDHA@) on chromosome
	NM 032935	

NM_003695	Homo sapiens lymphocyte antigen 6 complex, locus D (E48), mRNA
NM 006787	Homo sapiens melanoma antigen, family D, 2 (MAGED2), mRNA
NM 016205	Homo sapiens platelet derived growth factor C (PDGFC), mRNA
NM 017913	Homo sapiens Hsp90-associating relative of Cdc37 (HARC), mRNA
NM 017701	Homo sapiens Rho GTPase activating protein 8 (ARHGAP8), mRNA
NM 015366	Homo sapiens Rho GTPase activating protein 8 (ARHGAP8), mRNA
NM 012269	Homo sapiens hyaluronoglucosaminidase 4 (HYAL4), mRNA
NM 006207	Homo sapiens platelet-derived growth factor receptor-like (PDGFRL), mRNA
NM_004986	Homo sapiens kinectin 1 (kinesin receptor) (KTN1), mRNA
NM_001840	Homo sapiens cannabinoid receptor 1 (brain) (CNR1), transcript variant 1, mRNA
NM 014417	Homo sapiens Bcl-2 binding component 3 (BBC3), mRNA
NM_033223	Homo sapiens gamma-aminobutyric acid (GABA) A receptor, gamma 3 (GABRG3), mRNA
NM 005762	Homo sapiens tripartite motif-containing 28 (TRIM28), mRNA
NM_015906	Homo sapiens tripartite motif-containing 33 (TRIM33), transcript variant alpha, mRNA
NM_033020	Homo sapiens tripartite motif-containing 33 (TRIM33), transcript variant beta, mRNA
NM_032421	Homo sapiens cytoplasmic linker 2 (CYLN2), transcript variant 2, mRNA
NM_031416	Homo sapiens chromosome 18 open reading frame 2 (C18orf2), mRNA
NM_014412	Homo sapiens Siah-interacting protein (SIP), mRNA
NM_016212	Homo sapiens TP53TG3 protein (TP53TG3), mRNA
NM_016552	Homo sapiens testis specific ankyrin-like protein 1 (LOC51281), mRNA
NM_015369	Homo sapiens TP53TG3 protein (TP53TG3), mRNA
NM_033284	Homo sapiens transducin beta-like 1 protein (TBL1Y), mRNA
NM_031951	Homo sapiens NYD-SP11 protein (NYD-SP11), mRNA
NM_020414	Homo sapiens DEAD/H (Asp-Glu-Ala-Asp/His) box polypeptide 24 (DDX24), mRNA
NM_007268	Homo sapiens Ig superfamily protein (Z39IG), mRNA
NM_006707	Homo sapiens butyrophilin-like 3 (BTNL3), mRNA
NM_002491	Homo sapiens NADH dehydrogenase (ubiquinone) 1 beta subcomplex, 3 (12kD, B12) (NDUFB3), mRNA
NM_001386	Homo sapiens dihydropyrimidinase-like 2 (DPYSL2), mRNA
NM_000090	Homo sapiens collagen, type III, alpha 1 (Ehlers-Danlos syndrome type IV, autosomal dominant) (COL3A1), mRNA
NM_033150	Homo sapiens collagen, type II, alpha 1 (primary osteoarthritis, spondyloepiphyseal dysplasia, congenital) (COL2A1), transcript variant 2, mRNA
NM_001844	Homo sapiens collagen, type II, alpha 1 (primary osteoarthritis, spondyloepiphyseal dysplasia, congenital) (COL2A1), transcript variant 1, mRNA
NM_025245	Homo sapiens pre-B-cell leukemia transcription factor 4 (PBX4), mRNA
NM_004342	Homo sapiens caldesmon 1 (CALD1), transcript variant 3, mRNA
NM_033157	Homo sapiens caldesmon 1 (CALD1), transcript variant 2, mRNA
NM_033140	Homo sapiens caldesmon 1 (CALD1), transcript variant 5, mRNA
NM_033139	Homo sapiens caldesmon 1 (CALD1), transcript variant 4, mRNA
NM_033138	Homo sapiens caldesmon 1 (CALD1), transcript variant 1, mRNA
NM_032635	Homo sapiens seven transmembrane domain protein (NIFIE14), mRNA
NM_030912	Homo sapiens ring finger protein 27 (RNF27), mRNA
NM_019849	Homo sapiens solute carrier family 7, (cationic amino acid transporter, y+
	system) member 10 (SLC7A10), mRNA

NM 017844	Homo sapiens testis specific ankyrin-like protein 1 (LOC51281), mRNA
NM 014242	Homo sapiens zinc finger protein 237 (ZNF237), mRNA
NM 001715	Homo sapiens B lymphoid tyrosine kinase (BLK), mRNA
NM_033158	Homo sapiens hyaluronoglucosaminidase 2 (HYAL2), transcript variant 2,
	mRNA
NM_033159	Homo sapiens hyaluronoglucosaminidase 1 (HYAL1), transcript variant 2,
	mRNA
NM_007312	Homo sapiens hyaluronoglucosaminidase 1 (HYAL1), transcript variant 1,
	mRNA
NM_006119	Homo sapiens fibroblast growth factor 8 (androgen-induced) (FGF8), transcript
ND 6 022165	variant B, mRNA Homo sapiens fibroblast growth factor 8 (androgen-induced) (FGF8), transcript
NM_033165	variant A, mRNA
NM_033164	Homo sapiens fibroblast growth factor 8 (androgen-induced) (FGF8), transcript
14141_033104	variant E, mRNA
NM_033163	Homo sapiens fibroblast growth factor 8 (androgen-induced) (FGF8), transcript
1111_033103	variant F, mRNA
NM_002009	Homo sapiens fibroblast growth factor 7 (keratinocyte growth factor) (FGF7),
_	mRNA
NM_021907	Homo sapiens dystrobrevin, beta (DTNB), transcript variant 1, mRNA
NM_033148	Homo sapiens dystrobrevin, beta (DTNB), transcript variant 3, mRNA
NM_033147	Homo sapiens dystrobrevin, beta (DTNB), transcript variant 2, mRNA
NM_015902	Homo sapiens progestin induced protein (DD5), mRNA
NM_000777	Homo sapiens cytochrome P450, subfamily IIIA (niphedipine oxidase),
	polypeptide 5 (CYP3A5), mRNA
NM_000764	Homo sapiens cytochrome P450, subfamily IIA (phenobarbital-inducible),
17.6.0000	polypeptide 7 (CYP2A7), transcript variant 1, mRNA
NM_030589	Homo sapiens cytochrome P450, subfamily IIA (phenobarbital-inducible),
NM_000762	polypeptide 7 (CYP2A7), transcript variant 2, mRNA Homo sapiens cytochrome P450, subfamily IIA (phenobarbital-inducible),
NWI_000702	polypeptide 6 (CYP2A6), mRNA
NM 018957	Homo sapiens SH3-domain binding protein 1 (SH3BP1), mRNA
NM 033258	Homo sapiens G-protein gamma 8 subunit (GNG8), mRNA
NM 033260	Homo sapiens winged helix/forkhead transcription factor (HFH1), mRNA
NM 018476	Homo sapiens brain expressed, X-linked 1 (BEX1), mRNA
NM 022154	Homo sapiens up-regulated by BCG-CWS (LOC64116), mRNA
NM_003773	Homo sapiens hyaluronoglucosaminidase 2 (HYAL2), transcript variant 1,
	mRNA
NM_032794	Homo sapiens NG22 protein (NG22), mRNA
NM_030768	Homo sapiens integrin-linked kinase-associated serine/threonine phosphatase 2C
	(ILKAP), mRNA
NM_025257	Homo sapiens NG22 protein (NG22), mRNA
NM_020996	Homo sapiens fibroblast growth factor 6 (FGF6), mRNA
NM_016543	Homo sapiens sialic acid binding Ig-like lectin 7 (SIGLEC7), mRNA
NM_016134	Homo sapiens plasma glutamate carboxypeptidase (PGCP), mRNA
NM 014385	Homo sapiens sialic acid binding Ig-like lectin 7 (SIGLEC7), mRNA
NM 013287	Homo sapiens phosphoprotein enriched in astrocytes 15 (PEA15), mRNA
NM 006102	Homo sapiens plasma glutamate carboxypeptidase (PGCP), mRNA
NM 004112	Homo sapiens fibroblast growth factor 11 (FGF11), mRNA Homo sapiens fibroblast growth factor 10 (FGF10), mRNA
NM_004465 NM_003811	Homo sapiens tumor necrosis factor (ligand) superfamily, member 9 (TNFSF9),
14141_002011	mRNA
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NM 003063	Homo sapiens sarcolipin (SLN), mRNA
NM 003768	Homo sapiens phosphoprotein enriched in astrocytes 15 (PEA15), mRNA
NM 002010	Homo sapiens fibroblast growth factor 9 (glia-activating factor) (FGF9), mRNA
NM 033215	Homo sapiens protein phosphatase 1, regulatory (inhibitor) subunit 3F
	(PPP1R3F), mRNA
NM 032741	Homo sapiens 1-acylglycerol-3-phosphate O-acyltransferase 1 (lysophosphatidic
	acid acyltransferase, alpha) (AGPAT1), mRNA
NM 022152	Homo sapiens PP1201 protein (PP1201), mRNA
NM 033225	Homo sapiens CUB and Sushi multiple domains 1 (CSMD1), mRNA
NM 014505	Homo sapiens potassium large conductance calcium-activated channel,
1111_014505	subfamily M, beta member 4 (KCNMB4), mRNA
NM 032559	Homo sapiens kinesin protein (LOC84643), mRNA
NM 015394	Homo sapiens zinc finger protein 10 (KOX 1) (ZNF10), mRNA
NM 003388	Homo sapiens cytoplasmic linker 2 (CYLN2), transcript variant 1, mRNA
NM 032736	Homo sapiens torsin family 1, member B (torsin B) (TOR1B), mRNA
NM 032689	Homo sapiens hypothetical protein MGC13071 (MGC13071), mRNA
NM 032227	Homo sapiens hypothetical protein FLJ22679 (FLJ22679), mRNA
NM_014506	Homo sapiens torsin family 1, member B (torsin B) (TOR1B), mRNA
NM 030900	Homo sapiens cell cycle progression 2 protein (CPR2), mRNA
NM 030758	Homo sapiens oxysterol binding protein 2 (OSBP2), mRNA
NM 017698	Homo sapiens bypothetical protein FLJ22679 (FLJ22679), mRNA
NM 018225	Homo sapiens homolog of C. elegans smu-1 (SMU-1), mRNA
NM 016285	Homo sapiens Kruppel-like factor 12 (KLF12), mRNA
	Homo sapiens Kruppel-like factor 12 (KLF12), mRNA
NM_007249	Homo sapiens kruppei-like factor 12 (KLF12), filktva Homo sapiens trans-golgi network protein 2 (TGOLN2), mRNA
NM_006464	Homo sapiens trans-goigi network protein 2 (100EN2), mixty. Homo sapiens 1-acylglycerol-3-phosphate O-acyltransferase 1 (lysophosphatidic
NM_006411	
NR 6 004740	acid acyltransferase, alpha) (AGPAT1), mRNA Homo sapiens cell cycle progression 2 protein (CPR2), mRNA
NM_004749	Homo sapiens cen cycle progression 2 protein (CFR2), inktyA Homo sapiens peptidase D (PEPD), mRNA
NM_000285	Homo sapiens glucose-6-phosphatase, transport (glucose-6-phosphate) protein 1
NM_001467	(G6PT1), mRNA
NM 033198	Homo sapiens phosphatidylinositol glycan, class S (PIGS), mRNA
NM 002920	Homo sapiens regulatory factor X, 4 (influences HLA class II expression)
-	(RFX4), mRNA
NM_018944	Homo sapiens chromosome 21 open reading frame 45 (C21orf45), mRNA
NM 033214	Homo sapiens glycerol kinase pseudogene 2 (GKP2), mRNA
NM_033089	Homo sapiens hypothetical protein FLJ22115 (FLJ22115), mRNA
NM_016015	Homo sapiens leucine carboxyl methyltransferase (LCMT), mRNA
NM_033209	Homo sapiens Thy-1 co-transcribed (LOC94105), mRNA
NM_033093	Homo sapiens tripartite motif-containing 5 (TRIM5), transcript variant delta,
ND 4 022222	mRNA
NM_033092	Homo sapiens tripartite motif-containing 5 (TRIM5), transcript variant gamma, mRNA
NM 033091	Homo sapiens tripartite motif-containing 4 (TRIM4), transcript variant beta,
	mRNA
NM 033017	Homo sapiens tripartite motif-containing 4 (TRIM4), transcript variant alpha,
	mRNA
NM 033034	Homo sapiens tripartite motif-containing 5 (TRIM5), transcript variant alpha,
	mRNA
NM 015318	Homo sapiens Rho-specific guanine nucleotide exchange factor p114 (P114-
	RHO-GEF), mRNA
NM 007204	Homo sapiens DEAD/H (Asp-Glu-Ala-Asp/His) box polypeptide 20, 103kD

	(DDX20), mRNA
NM 032864	Homo sapiens hypothetical protein FLJ14936 (FLJ14936), mRNA
NM_032639	Homo sapiens phosphoinositol 4-phosphate adaptor protein-2 (FAPP2), mRNA
NM 032583	Homo sapiens ATP-binding cassette, sub-family C (CFTR/MRP), member 11
14141_032363	(ABCC11), mRNA
NM_032284	Homo sapiens hypothetical protein FLJ14936 (FLJ14936), mRNA
NM 032182	Homo sapiens hypothetical protein FLJ13614 (FLJ13614), mRNA
NM_021727	Homo sapiens fatty acid desaturase 3 (FADS3), mRNA
NM 022726	Homo sapiens elongation of very long chain fatty acids (FEN1/Elo2, SUR4/Elo3,
	yeast)-like 4 (ELOVL4), mRNA
NM_015162	Homo sapiens lipidosin (BG1), mRNA
NM_021176	Homo sapiens islet-specific glucose-6-phosphatase catalytic subunit-related protein (IGRP), mRNA
NM_019094	Homo sapiens nudix (nucleoside diphosphate linked moiety X)-type motif 4 (NUDT4), mRNA
NM 019091	Homo sapiens pleckstrin homology domain-containing, family A
14141_019091	(phosphoinositide binding specific) member 3 (PLEKHA3), mRNA
NM 018293	Homo sapiens hypothetical protein FLJ10997 (FLJ10997), mRNA
NM_015994	Homo sapiens ATPase, H+ transporting lysosomal (vacuolar proton pump), member M (ATP6M), mRNA
NM 015952	Homo sapiens PTD013 protein (PTD013), mRNA
NM 015899	Homo sapiens putative glycolipid transfer protein (LOC51054), mRNA
NM 016309	Homo sapiens leucine carboxyl methyltransferase (LCMT), mRNA
NM 013345	Homo sapiens G protein-coupled receptor (G2A), mRNA
NM 012228	Homo sapiens pilin-like transcription factor (PILB), mRNA
NM 006886	Homo sapiens ATP synthase, H+ transporting, mitochondrial F1 complex,
	epsilon subunit (ATP5E), mRNA
NM 002200	Homo sapiens interferon regulatory factor 5 (IRF5), transcript variant 1, mRNA
NM 032643	Homo sapiens interferon regulatory factor 5 (IRF5), transcript variant 2, mRNA
NM 004464	Homo sapiens fibroblast growth factor 5 (FGF5), transcript variant 1, mRNA
NM 033143	Homo sapiens fibroblast growth factor 5 (FGF5), transcript variant 2, mRNA
NM_020638	Homo sapiens fibroblast growth factor 23 (FGF23), mRNA
NM_000800	Homo sapiens fibroblast growth factor 1 (acidic) (FGF1), transcript variant 1,
	mRNA
NM_033137	Homo sapiens fibroblast growth factor 1 (acidic) (FGF1), transcript variant 3,
	mRNA
NM_032102	Homo sapiens Splicing factor, arginine/serine-rich, 46kD (SRP46), mRNA
NM_033136	Homo sapiens fibroblast growth factor 1 (acidic) (FGF1), transcript variant 2, mRNA
NM_002952	Homo sapiens ribosomal protein S2 (RPS2), mRNA
NM_033130	Homo sapiens sialic acid binding Ig-like lectin 10 (SIGLEC10), mRNA
NM_020665	Homo sapiens kidney-specific membrane protein (NX-17), mRNA
NM_033180	Homo sapiens olfactory receptor, family 51, subfamily B, member 2 (OR51B2), mRNA
NM_033179	Homo sapiens olfactory receptor, family 51, subfamily B, member 4 (OR51B4),
NIM 022170	mRNA
NM_033178	Homo sapiens double homeobox, 4 (DUX4), mRNA
NM_033049	Homo sapiens mucin 13, epithelial transmembrane (MUC13), mRNA
NM_021619	Homo sapiens PR domain containing 12 (PRDM12), mRNA
NM 020382	Homo sapiens PR/SET domain containing protein 07 (SET07), mRNA
NM 007365	Homo sapiens peptidyl arginine deiminase, type II (PDI2), mRNA
NM_015894	Homo sapiens stathmin-like 3 (STMN3), mRNA

NM 032491	Homo sapiens regulatory factor X, 4 (influences HLA class II expression)
NM_032491	(RFX4), mRNA
NM_024551	Homo sapiens hypothetical protein FLJ21432 (FLJ21432), mRNA
NM_021830	Homo sapiens chromosome 10 open reading frame 2 (C10orf2), mRNA
NM_017972	Homo sapiens hypothetical protein FLJ20689 (FLJ20689), mRNA
NM_020398	Homo sapiens serine protease inhibitor-like, with Kunitz and WAP domains 1 (eppin) (SPINLW1), mRNA
NM 020637	Homo sapiens fibroblast growth factor 22 (FGF22), mRNA
NM 019113	Homo sapiens fibroblast growth factor 21 (FGF21), mRNA
NM 017926	Homo sapiens hypothetical protein FLJ20689 (FLJ20689), mRNA
NM 016444	Homo sapiens zinc finger protein 226 (ZNF226), mRNA
NM 015966	Homo sapiens serologically defined breast cancer antigen 84 (SDBCAG84),
	mRNA
NM 015919	Homo sapiens zinc finger protein 226 (ZNF226), mRNA
NM 015474	Homo sapiens SAM domain and HD domain, 1 (SAMHD1), mRNA
NM_007096	Homo sapiens clathrin, light polypeptide (Lca) (CLTA), transcript variant brain- specific, mRNA
NM_002007	Homo sapiens fibroblast growth factor 4 (heparin secretory transforming protein 1, Kaposi sarcoma oncogene) (FGF4), mRNA
NM_001833	Homo sapiens clathrin, light polypeptide (Lca) (CLTA), transcript variant
	nonbrain, mRNA
NM_022143	Homo sapiens NAG14 protein (NAG14), mRNA
NM_005292	Homo sapiens G protein-coupled receptor 18 (GPR18), mRNA
NM_001371	Homo sapiens dynein, axonemal, heavy polypeptide 8 (DNAH8), mRNA
NM_012276	Homo sapiens leukocyte immunoglobulin-like receptor, subfamily A (without TM domain), member 4 (ILT7), mRNA
NM_012092	Homo sapiens inducible T-cell co-stimulator (ICOS), mRNA
NM_032447	Homo sapiens fibrillin3 (KIAA1776), mRNA
NM_024017	Homo sapiens homeo box B9 (HOXB9), mRNA
NM_019558	Homo sapiens homeo box D8 (HOXD8), mRNA
NM_032379	Homo sapiens synaptotagmin-like 2 (SYTL2), transcript variant b, mRNA
NM_024690	Homo sapiens mucin 16 (MUC16), mRNA
NM_018558	Homo sapiens gamma-aminobutyric acid (GABA) receptor, theta (GABRQ), mRNA
NM_014452	Homo sapiens tumor necrosis factor receptor superfamily, member 21 (TNFRSF21), mRNA
NM_006242	Homo sapiens protein phosphatase 1, regulatory subunit 3D (PPP1R3D), mRNA
NM_006545	Homo sapiens homologous to yeast nitrogen permease (candidate tumor suppressor) (NPR2L), mRNA
NM_005398	Homo sapiens protein phosphatase 1, regulatory (inhibitor) subunit 3C (PPP1R3C), mRNA
NM_006645	Homo sapiens serologically defined colon cancer antigen 28 (SDCCAG28), mRNA
NM 032800	Homo sapiens hypothetical protein FLJ14525 (FLJ14525), mRNA
NM 004265	
NM_013402	
NM_031428	
NM 025243	
NM_024411	
NM_007368	Homo sapiens RAS p21 protein activator (GTPase activating protein) 3
NM 003912	
NM 013402 NM 031428 NM 025243 NM 024411	Homo sapiens fatty acid desaturase 2 (FADS2), mRNA Homo sapiens fatty acid desaturase 1 (FADS1), mRNA Homo sapiens hypothetical protein FLJ14525 (FLJ14525), mRNA Homo sapiens solute carrier family 19, member 3 (SLC19A3), mRNA Homo sapiens prodynorphin (PDYN), mRNA

NTM 015004	Homo sapiens ubiquitin C-terminal hydrolase UCH37 (UCH37), mRNA
NM_015984	Homo sapiens angiopoietin-like 4 (ANGPTL4), mRNA
NM_016109	Homo sapiens myotubularin related protein 2 (MTMR2), mRNA
NM_016156	Homo sapiens progesterone receptor membrane component 1 (PGRMC1),
NM_006667	mRNA
NM 006312	Homo sapiens nuclear receptor co-repressor 2 (NCOR2), mRNA
NM_006320	Homo sapiens progesterone receptor membrane component 2 (PGRMC2), mRNA
NM 000441	Homo sapiens solute carrier family 26, member 4 (SLC26A4), mRNA
NM 032995	Homo sapiens Rho guanine nucleotide exchange factor (GEF) 4 (ARHGEF4),
	transcript variant 2, mRNA
NM 015320	Homo sapiens Rho guanine nucleotide exchange factor (GEF) 4 (ARHGEF4),
1111_010020	transcript variant 1, mRNA
NM 014448	Homo sapiens Rho guanine exchange factor (GEF) 16 (ARHGEF16), mRNA
NM 005435	Homo sapiens Rho guanine nucleotide exchange factor (GEF) 5 (ARHGEF5),
1447_000 155	mRNA
NM 004723	Homo sapiens rho/rac guanine nucleotide exchange factor (GEF) 2 (ARHGEF2),
144_001725	mRNA
NM_004706	Homo sapiens Rho guanine nucleotide exchange factor (GEF) 1 (ARHGEF1),
111/1_00 1700	mRNA
NM 001031	Homo sapiens ribosomal protein S28 (RPS28), mRNA
NM 001030	Homo sapiens ribosomal protein S27 (metallopanstimulin 1) (RPS27), mRNA
NM 001029	Homo sapiens ribosomal protein S26 (RPS26), mRNA
NM_002913	Homo sapiens replication factor C (activator 1) 1 (145kD) (RFC1), mRNA
NM 005685	Homo sapiens GTF2I repeat domain-containing 1 (GTF2IRD1), transcript
14141_005085	variant 2, mRNA
NM_005117	Homo sapiens fibroblast growth factor 19 (FGF19), mRNA
NM 001363	Homo sapiens dyskeratosis congenita 1, dyskerin (DKC1), mRNA
NM_005765	Homo sapiens ATPase, H+ transporting, lysosomal (vacuolar proton pump)
27.5 001040	membrane sector associated protein M8-9 (APT6M8-9), mRNA
NM_001848	Homo sapiens collagen, type VI, alpha 1 (COL6A1), mRNA
NM_004932	Homo sapiens cadherin 6, type 2, K-cadherin (fetal kidney) (CDH6), mRNA
NM_005673	Homo sapiens solute carrier family 25 (mitochondrial carrier; Graves disease
	autoantigen), member 16 (SLC25A16), nuclear gene encoding mitochondrial
	protein, mRNA
NM_032943	Homo sapiens synaptotagmin-like 2 (SYTL2), transcript variant a, mRNA
NM_006932	Homo sapiens smoothelin (SMTN), mRNA
NM_000411	Homo sapiens holocarboxylase synthetase (biotin-[proprionyl-Coenzyme A-
<u> </u>	carboxylase (ATP-hydrolysing)] ligase) (HLCS), mRNA
NM_030777	Homo sapiens solute carrier family 2 (facilitated glucose transporter), member 10
	(SLC2A10), mRNA
NM_022897	Homo sapiens RAN binding protein 17 (RANBP17), mRNA
NM_015339	Homo sapiens activity-dependent neuroprotector (ADNP), mRNA
NM_015024	Homo sapiens RAN binding protein 16 (RANBP16), mRNA
NM_022046	Homo sapiens kallikrein 14 (KLK14), mRNA
NM_020041	Homo sapiens solute carrier family 2 (facilitated glucose transporter), member 9 (SLC2A9), mRNA
NM 019851	Homo sapiens fibroblast growth factor 20 (FGF20), mRNA
NM 019555	Homo sapiens Rho guanine nucleotide exchange factor (GEF) 3 (ARHGEF3),
	mRNA
NM_016277	Homo sapiens RAB23, member RAS oncogene family (RAB23), mRNA
NM 014629	Homo sapiens Rho guanine nucleotide exchange factor (GEF) 10 (ARHGEF10),
	1 (ODI) 10 (Maiori 19)

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ND (00(000	mRNA
NM_006989	Homo sapiens Ca2+-promoted Ras inactivator (CAPRI), mRNA
NM_006568	Homo sapiens cell growth regulatory with ring finger domain (CGR19), mRNA
NM_004841	Homo sapiens RAS protein activator like 2 (RASAL2), mRNA
NM_004115	Homo sapiens fibroblast growth factor 14 (FGF14), mRNA
NM_003244	Homo sapiens TGFB-induced factor (TALE family homeobox) (TGIF), mRNA
NM_007285	Homo sapiens GABA(A) receptor-associated protein-like 2 (GABARAPL2), mRNA
NM_006047	Homo sapiens RNA binding motif protein 12 (RBM12), mRNA
NM_032588	Homo sapiens ring finger protein 28 (RNF28), mRNA
NM_030766	Homo sapiens apoptosis regulator BCL-G (BCLG), mRNA
NM_022788	Homo sapiens Purinergic receptor P2Y, G protein-coupled, 12 (P2RY12), mRNA
NM_015641	Homo sapiens testis derived transcript (3 LIM domains) (TES), mRNA
NM_018144	Homo sapiens Sec61 alpha form 2 (FLJ10578), mRNA
NM_032015	Homo sapiens ring finger protein 26 (RNF26), mRNA
NM_014713	Homo sapiens lysosomal-associated protein transmembrane 4 alpha
	(LAPTM4A), mRNA
NM_020415	Homo sapiens found in inflammatory zone 3 (FIZZ3), mRNA
NM_020358	Homo sapiens ring finger protein 18 (RNF18), mRNA
NM_005882	Homo sapiens macrophage erythroblast attacher (MAEA), mRNA
NM_016523	Homo sapiens killer cell lectin-like receptor subfamily F, member 1 (KLRF1), mRNA
NM_014141	Homo sapiens contactin associated protein-like 2 (CNTNAP2), mRNA
NM_006862	Homo sapiens tudor and KH domain-containing protein (TDRKH), mRNA
NM_006779	Homo sapiens Cdc42 effector protein 2 (CEP2), mRNA
NM_006292	Homo sapiens tumor susceptibility gene 101 (TSG101), mRNA
NM_006449	Homo sapiens Cdc42 effector protein 3 (CEP3), mRNA
NM_002558	Homo sapiens purinergic receptor P2X, ligand-gated ion channel, 1 (P2RX1), mRNA
NM_006712	Homo sapiens FAST kinase (FASTK), transcript variant 1, mRNA
NM_033015	Homo sapiens FAST kinase (FASTK), transcript variant 2, mRNA
NM_025096	Homo sapiens FAST kinase (FASTK), transcript variant 3, mRNA
NM_003852	Homo sapiens transcriptional intermediary factor 1 (TIF1), mRNA
NM_003770	Homo sapiens keratin, hair, acidic, 7 (KRTHA7), mRNA
NM_021013	Homo sapiens keratin, hair, acidic, 4 (KRTHA4), mRNA
NM_004068	Homo sapiens adaptor-related protein complex 2, mu 1 subunit (AP2M1), mRNA
NM_006803	Homo sapiens adaptor-related protein complex 3, mu 2 subunit (AP3M2), mRNA
NM_005498	Homo sapiens adaptor-related protein complex 1, mu 2 subunit (AP1M2), mRNA
NM_032981	Homo sapiens dystrobrevin, alpha (DTNA), transcript variant zeta, mRNA
NM_032980	Homo sapiens dystrobrevin, alpha (DTNA), transcript variant epsilon, mRNA
NM_032979	Homo sapiens dystrobrevin, alpha (DTNA), transcript variant gamma, mRNA
NM_032978	Homo sapiens dystrobrevin, alpha (DTNA), transcript variant beta, mRNA
NM_032975	Homo sapiens dystrobrevin, alpha (DTNA), transcript variant alpha, mRNA
NM_001392	Homo sapiens dystrobrevin, alpha (DTNA), transcript variant DTN3, mRNA
NM_001391	Homo sapiens dystrobrevin, alpha (DTNA), transcript variant DTN2, mRNA
NM_001390	Homo sapiens dystrobrevin, alpha (DTNA), transcript variant DTN1, mRNA
NM_001026	Homo sapiens ribosomal protein S24 (RPS24), transcript variant 2, mRNA
NM 033022	Homo sapiens ribosomal protein S24 (RPS24), transcript variant 1, mRNA
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NM_023014 Homo sapiens osteoglycin (osteoinductive factor, mimecan) (OGN), transcript variant 2, mRNA		
NM_013014	NM_024416	
NM_016152 Homo sapiens osteoglycin (osteoinductive factor, mimecan) (OGN), transcript variant 3, mRNA NM_016152 Homo sapiens retinoic acid receptor, beta (RARB), transcript variant 2, mRNA NM_032971 Homo sapiens retinoic acid receptor, beta (RARB), transcript variant 1, mRNA NM_032972 Homo sapiens caspase 10, apoptosis-related cysteine protease (CASP10), transcript variant D, mRNA Homo sapiens caspase 10, apoptosis-related cysteine protease (CASP10), transcript variant D, mRNA Homo sapiens caspase 10, apoptosis-related cysteine protease (CASP10), transcript variant B, mRNA Homo sapiens caspase 10, apoptosis-related cysteine protease (CASP10), transcript variant B, mRNA Homo sapiens caspase 10, apoptosis-related cysteine protease (CASP6), transcript variant tea, mRNA Homo sapiens caspase 6, apoptosis-related cysteine protease (CASP6), transcript variant alpha, mRNA Homo sapiens caspase 6, apoptosis-related cysteine protease (CASP6), transcript variant alpha, mRNA Homo sapiens 2;3'-cyclic nucleotide 3' phosphodiesterase (CNP), mRNA NM_033133 Homo sapiens 2;3'-cyclic nucleotide 3' phosphodiesterase (CNP), mRNA Homo sapiens syrapanic cation transporter OKB1 (OKB1), mRNA Homo sapiens syrapanic cation transporter OKB1 (OKB1), mRNA Homo sapiens syrataxin 18 (STX18), mRNA Homo sapiens syrataxin 18 (STX18), mRNA Homo sapiens syrataxin 18 (STX18), mRNA Homo sapiens syntaxin 18 (STX18), mRNA Homo sapiens shosphodiesterase 3B, CBLA3A), mRNA Homo sapiens shosphodiesterase 3B, cGMP-inhibited (PDE3B), mRNA Homo sapiens shosphodiesterase 3B, cGMP-inhibited (PDE3B), mRNA Homo sapiens shosphodiesterase 3B, cGMP-inhibited (PDE3B), mRNA Homo sapiens scaspase 2, apoptosis-related cysteine protein (SHARP), mRNA Homo sapiens caspase 2, apoptosis-related cysteine protease (neural precursor cell expressed, development	NM_033014	Homo sapiens osteoglycin (osteoinductive factor, mimecan) (OGN), transcript
NM 0032974 Homo sapiens retinoic acid receptor, beta (RARB), transcript variant 1, mRNA NM 032976 Homo sapiens caspase 10, apoptosis-related cysteine protease (CASP10), transcript variant D, mRNA NM_032976 Homo sapiens caspase 10, apoptosis-related cysteine protease (CASP10), transcript variant C, mRNA NM_032974 Homo sapiens caspase 10, apoptosis-related cysteine protease (CASP10), transcript variant B, mRNA NM_001230 Homo sapiens caspase 10, apoptosis-related cysteine protease (CASP10), transcript variant B, mRNA NM_001230 Homo sapiens caspase 10, apoptosis-related cysteine protease (CASP10), transcript variant than, mRNA NM_001296 Homo sapiens caspase 6, apoptosis-related cysteine protease (CASP6), transcript variant beta, mRNA NM_001226 Homo sapiens caspase 6, apoptosis-related cysteine protease (CASP6), transcript variant alpha, mRNA NM_033133 Homo sapiens 2;3'-cyclic nucleotide 3' phosphodiesterase (CNP), mRNA NM_033125 Homo sapiens organic cation transporter OKB1 (OKB1), mRNA NM_000349 Homo sapiens ankyrin repeat domain 2 (stretch responsive muscle) (ANKRD2), mRNA NM_000540 Homo sapiens syntaxin 18 (STX18), mRNA NM_014808 Homo sapiens KIAA0793 gene product (KIAA0793), mRNA NM_005428 Homo sapiens kIAA0793 gene product (KIAA0793), mRNA NM_005428 Homo sapiens base 3A, pancreatic (protease E) (ELA3A), mRNA NM_005427 Homo sapiens phosphodiesterase 3B, cGMP-inhibited (PDE3B), mRNA NM_003069 Homo sapiens shand adult testis expressed transcript protein (FATE), mRNA NM_033069 Homo sapiens SMART/HDAC1 associated repressor protein (SHARP), mRNA NM_032984 Homo sapiens SMART/HDAC1 associated repressor protein (SHARP), mRNA NM_032985 Homo sapiens fetal and adult testis expressed transcript protein (FATE), mRNA NM_032986 Homo sapiens facapase 2, apoptosis-related cysteine protease (neural precursor cell expressed, developmentally down-regulated 2) (CASP2), transcript variant 1, mRNA NM_032987 Homo sapiens tumor necrosis factor receptor superfamily, member 6b, decoy (TNFRSF6B), transcript variant 1, mRNA NM_032985 Homo s	NM_014057	Homo sapiens osteoglycin (osteoinductive factor, mimecan) (OGN), transcript
NM 0032974 Homo sapiens retinoic acid receptor, beta (RARB), transcript variant 1, mRNA NM 032976 Homo sapiens caspase 10, apoptosis-related cysteine protease (CASP10), transcript variant D, mRNA NM_032976 Homo sapiens caspase 10, apoptosis-related cysteine protease (CASP10), transcript variant C, mRNA NM_032974 Homo sapiens caspase 10, apoptosis-related cysteine protease (CASP10), transcript variant B, mRNA NM_001230 Homo sapiens caspase 10, apoptosis-related cysteine protease (CASP10), transcript variant B, mRNA NM_001230 Homo sapiens caspase 10, apoptosis-related cysteine protease (CASP10), transcript variant than, mRNA NM_001296 Homo sapiens caspase 6, apoptosis-related cysteine protease (CASP6), transcript variant beta, mRNA NM_001226 Homo sapiens caspase 6, apoptosis-related cysteine protease (CASP6), transcript variant alpha, mRNA NM_033133 Homo sapiens 2;3'-cyclic nucleotide 3' phosphodiesterase (CNP), mRNA NM_033125 Homo sapiens organic cation transporter OKB1 (OKB1), mRNA NM_000349 Homo sapiens ankyrin repeat domain 2 (stretch responsive muscle) (ANKRD2), mRNA NM_000540 Homo sapiens syntaxin 18 (STX18), mRNA NM_014808 Homo sapiens KIAA0793 gene product (KIAA0793), mRNA NM_005428 Homo sapiens kIAA0793 gene product (KIAA0793), mRNA NM_005428 Homo sapiens base 3A, pancreatic (protease E) (ELA3A), mRNA NM_005427 Homo sapiens phosphodiesterase 3B, cGMP-inhibited (PDE3B), mRNA NM_003069 Homo sapiens shand adult testis expressed transcript protein (FATE), mRNA NM_033069 Homo sapiens SMART/HDAC1 associated repressor protein (SHARP), mRNA NM_032984 Homo sapiens SMART/HDAC1 associated repressor protein (SHARP), mRNA NM_032985 Homo sapiens fetal and adult testis expressed transcript protein (FATE), mRNA NM_032986 Homo sapiens facapase 2, apoptosis-related cysteine protease (neural precursor cell expressed, developmentally down-regulated 2) (CASP2), transcript variant 1, mRNA NM_032987 Homo sapiens tumor necrosis factor receptor superfamily, member 6b, decoy (TNFRSF6B), transcript variant 1, mRNA NM_032985 Homo s	NM 016152	Homo sapiens retinoic acid receptor, beta (RARB), transcript variant 2, mRNA
NM_032977 Homo sapiens caspase 10, apoptosis-related cysteine protease (CASP10), transcript variant D, mRNA	NM 000965	
transcript variant C, mRNA NM_032974 Homo sapiens caspase 10, apoptosis-related cysteine protease (CASP10), transcript variant B, mRNA NM_001230 Homo sapiens caspase 10, apoptosis-related cysteine protease (CASP10), transcript variant A, mRNA NM_032992 Homo sapiens caspase 6, apoptosis-related cysteine protease (CASP6), transcript variant beta, mRNA NM_001226 Homo sapiens caspase 6, apoptosis-related cysteine protease (CASP6), transcript variant alpha, mRNA NM_033133 Homo sapiens organic cation transporter OKB1 (OKB1), mRNA NM_033125 Homo sapiens organic cation transporter OKB1 (OKB1), mRNA NM_033125 Homo sapiens ankyrin repeat domain 2 (stretch responsive muscle) (ANKRD2), mRNA NM_000540 Homo sapiens ryanodine receptor 1 (skeletal) (RYR1), mRNA NM_016930 Homo sapiens syntaxin 18 (STX18), mRNA NM_014808 Homo sapiens KIAA0793 gene product (KIAA0793), mRNA NM_005747 Homo sapiens elastase 3A, pancreatic (protease E) (ELA3A), mRNA NM_005747 Homo sapiens phosphodiesterase 3B, GGMP-inhibited (PDE3B), mRNA NM_033069 Homo sapiens SMART/HDAC1 associated repressor protein (FATE), mRNA NM_033085 Homo sapiens SMART/HDAC1 associated repressor protein (SHARP), mRNA NM_032984 Homo sapiens caspase 2, apoptosis-related cysteine protease (neural precursor cell expressed, developmentally down-regulated 2) (CASP2), transcript variant 4, mRNA NM_032983 Homo sapiens caspase 2, apoptosis-related cysteine protease (neural precursor cell expressed, developmentally down-regulated 2) (CASP2), transcript variant 1, mRNA NM_032983 Homo sapiens tumor necrosis factor receptor superfamily, member 6b, decoy (TNFRSF6B), transcript variant 1, mRNA NM_032984 Homo sapiens tumor necrosis factor receptor superfamily, member 6b, decoy (TNFRSF6B), transcript variant 1 (MRNA) NM_032985 Homo sapiens tumor necrosis factor receptor superfamily, member 6b, decoy (TNFRSF6B), transcript variant 1 (MRNA) NM_032985 Homo sapiens tumor necrosis factor receptor superfamily, member 6b, decoy (TNFRSF6B), transcript variant 3, mRNA	NM_032977	
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(TNFRSF6B), transcript variant 3, mRNA	NM_001224	Homo sapiens caspase 2, apoptosis-related cysteine protease (neural precursor cell expressed, developmentally down-regulated 2) (CASP2), transcript variant 2,
NM_033012 Homo sapiens tumor necrosis factor (ligand) superfamily, member 11		(TNFRSF6B), transcript variant 3, mRNA
	NM_033012	Homo sapiens tumor necrosis factor (ligand) superfamily, member 11

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27.5.00.701	(TNFSF11), transcript variant 2, mRNA
NM_003701	Homo sapiens tumor necrosis factor (ligand) superfamily, member 11
	(TNFSF11), transcript variant 1, mRNA
NM_005409	Homo sapiens small inducible cytokine subfamily B (Cys-X-Cys), member 11
3 D C 00 500 5	(SCYB11), mRNA
NM_005035	Homo sapiens polymerase (RNA) mitochondrial (DNA directed) (POLRMT),
	nuclear gene encoding mitochondrial protein, mRNA
NM_006980	Homo sapiens transcription termination factor, mitochondrial (MTERF), nuclear
	gene encoding mitochondrial protein, mRNA
NM_001305	Homo sapiens claudin 4 (CLDN4), mRNA
NM_032996	Homo sapiens caspase 9, apoptosis-related cysteine protease (CASP9), transcript
	variant beta, mRNA
NM_001229	Homo sapiens caspase 9, apoptosis-related cysteine protease (CASP9), transcript
	variant alpha, mRNA
NM_004346	Homo sapiens caspase 3, apoptosis-related cysteine protease (CASP3), transcript
	variant alpha, mRNA
NM_032991	Homo sapiens caspase 3, apoptosis-related cysteine protease (CASP3), transcript
	variant beta, mRNA
NM_033057	Homo sapiens olfactory receptor, family 2, subfamily B, member 2 (OR2B2),
	mRNA
NM_033051	Homo sapiens thymic stromal co-transporter (TSCOT), mRNA
NM_033048	Homo sapiens CPX chromosome region, candidate 1 (CPXCR1), mRNA
NM_033007	Homo sapiens death effector filament-forming Ced-4-like apoptosis protein
	(DEFCAP), transcript variant E, mRNA
NM_033006	Homo sapiens death effector filament-forming Ced-4-like apoptosis protein
	(DEFCAP), transcript variant D, mRNA
NM_033005	Homo sapiens death effector filament-forming Ced-4-like apoptosis protein
	(DEFCAP), transcript variant C, mRNA
NM_033004	Homo sapiens death effector filament-forming Ced-4-like apoptosis protein
	(DEFCAP), transcript variant A, mRNA
NM_014922	Homo sapiens death effector filament-forming Ced-4-like apoptosis protein
77.5.00000	(DEFCAP), transcript variant B, mRNA
NM_000088	Homo sapiens collagen, type I, alpha 1 (COL1A1), mRNA
NM_019105	Homo sapiens tenascin XB (TNXB), transcript variant XB, mRNA
NM_033036	Homo sapiens beta-galactose-3-O-sulfotransferase 3 (GAL3ST2), mRNA
NM_033029	Homo sapiens leishmanolysin-like (metallopeptidase M8 family) (LMLN),
\\\\\\\\\\\\\\\\\\\\\\\\\\\\\\\\\\\\\\	mRNA
NM_033028	Homo sapiens Bardet-Biedl syndrome 4 (BBS4), mRNA
NM_021807	Homo sapiens secretory protein SEC8 (SEC8), mRNA
NM_020137	Homo sapiens GRIP-associated protein 1 (GRASP1), mRNA
NM_015133	Homo sapiens mitogen-activated protein kinase 8 interacting protein 3
	(MAPK8IP3), mRNA
NM_014006	Homo sapiens PI-3-kinase-related kinase SMG-1 (SMG1), mRNA
NM_021914	Homo sapiens cofilin 2 (muscle) (CFL2), mRNA
NM_032520	Homo sapiens hypothetical protein CAB56184 (CAB56184), mRNA
NM_032923	Homo sapiens hypothetical protein MGC16025 (MGC16025), mRNA
NM_032917	Homo sapiens hypothetical protein MGC2848 (MGC2848), mRNA
NM_032868	Homo sapiens hypothetical protein FLJ14981 (FLJ14981), mRNA
NM_032862	Homo sapiens hypothetical protein FLJ14926 (FLJ14926), mRNA
NM_032801	Homo sapiens hypothetical protein FLJ14529 (FLJ14529), mRNA
NM_032753	Homo sapiens hypothetical protein MGC15631 (MGC15631), mRNA
NM_032737	Homo sapiens hypothetical protein MGC2721 (MGC2721), mRNA

NM_032668	Homo sapiens hypothetical protein MGC4771 (MGC4771), mRNA
NM_032503	Homo sapiens G protein-coupled receptor slt (SLT), mRNA
NM_032377	Homo sapiens hypothetical prot in MGC4549 (MGC4549), mRNA
NM 032326	Homo sapiens hypothetical protein MGC4618 (MGC4618), mRNA
NM 032306	Homo sapiens hypothetical protein MGC10974 (MGC10974), mRNA
NM 032281	Homo sapiens hypothetical protein DKFZp547J036 (DKFZp547J036), mRNA
NM 015650	Homo sapiens microtubule-interacting protein that associates with TRAF3 (MIP-
	T3), mRNA
NM 031487	Homo sapiens hypothetical protein MGC4604 (MGC4604), mRNA
NM 031470	Homo sapiens junctional adhesion molecule 3 (JAM3), mRNA
NM 031304	Homo sapiens hypothetical protein MGC4293 (MGC4293), mRNA
NM 031213	Homo sapiens hypothetical protein MGC:5244, (MGC:5244), mRNA
NM 031208	Homo sapiens hypothetical protein DKFZp566J2046 (DKFZP566J2046), mRNA
NM 030924	Homo sapiens hypothetical protein PRTD-NY3 (PRTD-NY3), mRNA
NM 030824	Homo sapiens hypothetical protein FLJ14356 (FLJ14356), mRNA
NM 030631	Homo sapiens solute carrier family 25 (mitochondrial oxodicarboxylate carrier),
11111_050051	member 21 (SLC25A21), mRNA
NM 024571	Homo sapiens hypothetical protein FLJ22940 (FLJ22940), mRNA
NM 025015	Homo sapiens KIAA0417 gene product (KIAA0417), mRNA
NM 024103	Homo sapiens hypothetical protein MGC2615 (MGC2615), mRNA
NM 030578	Homo sapiens hypothetical protein MGC4093 (MGC4093), mRNA
NM 014015	Homo sapiens MYLE protein (MYLE), mRNA
NM 025094	Homo sapiens hypothetical protein FLJ22184 (FLJ22184), mRNA
NM 025078	Homo sapiens hypothetical protein FLJ22378 (FLJ22378), mRNA
NM 025061	Homo sapiens hypothetical protein FLJ23420 (FLJ23420), mRNA
NM 024967	
NM 024898	Homo sapiens hypothetical protein FLJ11637 (FLJ11637), mRNA
	Homo sapiens hypothetical protein FLJ22757 (FLJ22757), mRNA
NM_024877	Homo sapiens hypothetical protein FLJ13265 (FLJ13265), mRNA
NM_024726	Homo sapiens hypothetical protein FLJ22527 (FLJ22527), mRNA
NM_024719	Homo sapiens hypothetical protein FLJ22474 (FLJ22474), mRNA
NM_024600	Homo sapiens hypothetical protein FLJ20898 (FLJ20898), mRNA
NM_024508	Homo sapiens hypothetical protein MGC10796 (MGC10796), mRNA
NM_024341	Homo sapiens hypothetical protein MGC4054 (MGC4054), mRNA
NM_024064	Homo sapiens hypothetical protein MGC5363 (MGC5363), mRNA
NM_024029	Homo sapiens hypothetical protein MGC3262 (MGC3262), mRNA
NM_023078	Homo sapiens hypothetical protein FLJ13852 (FLJ13852), mRNA
NM_023076	Homo sapiens hypothetical protein FLJ23360 (FLJ23360), mRNA
NM_022842	Homo sapiens hypothetical protein FLJ22969 (FLJ22969), mRNA
NM_022737	Homo sapiens hypothetical protein FLJ13055 (FLJ13055), mRNA
NM_022459	Homo sapiens hypothetical protein FLJ13046 similar to exportin 4; KIAA1721
17.6.655.65	pr (FLJ13046), mRNA
NM_022437	Homo sapiens ATP-binding cassette, sub-family G (WHITE), member 8 (sterolin
	2) (ABCG8), mRNA
NM_022135	Homo sapiens popeye protein 2 (POP2), mRNA
NM_022066	Homo sapiens likely ortholog of mouse ubiquitin-conjugating enzyme E2-230K (E2-230K), mRNA
NM_015480	Homo sapiens nectin 3 (DKFZP566B0846), mRNA
NM_004240	Homo sapiens thyroid hormone receptor interactor 10 (TRIP10), mRNA
NM 003589	Homo sapiens cullin 4A (CUL4A), mRNA
NM 021731	Homo sapiens hypothetical protein PP3501 (PP3501), mRNA
NM 020129	Homo sapiens placental protein 13-like protein (LOC56891), mRNA
NM 020196	Homo sapiens HCNP protein; XPA-binding protein 2 (HCNP), mRNA
11.12 020170	sapiens item process, At 11-0 moing process 2 (HCMF), like MA

NM_020224	Homo sapiens hypothetical protein DKFZp547O146 (DKFZp547O146), mRNA
NM_019064	Homo sapiens hypothetical protein (FLJ10832), mRNA
NM_019012	Homo sapiens phosphoinositol 3-phosphate-binding protein-2 (PEPP2), mRNA
NM_018635	Homo sapiens hypothetical protein PRO2900 (PRO2900), mRNA
NM_018687	Homo sapiens hepatocellular carcinoma-associated gene TD26 (LOC55908), mRNA
NM_018441	Homo sapiens peroxisomal trans 2-enoyl CoA reductase; putative short chain alcohol dehydrogenase (HSA250303), mRNA
NM 018645	Homo sapiens hypothetical protein HES6 (HES6), mRNA
NM 017967	Homo sapiens hypothetical protein FLJ20850 (FLJ20850), mRNA
NM 017914	Homo sapiens hypothetical protein FLJ20640 (FLJ20640), mRNA
NM 017905	Homo sapiens hypothetical protein FLJ20623 (FLJ20623), mRNA
NM 017722	Homo sapiens hypothetical protein FLJ20244 (FLJ20244), mRNA
NM_017668	Homo sapiens LIS1-interacting protein NUDE1, rat homolog (NUDE1), mRNA
NM 017616	Homo sapiens hypothetical protein FLJ20004 (FLJ20004), mRNA
NM 018185	Homo sapiens hypothetical protein FLJ10704 (FLJ10704), mRNA
NM 018074	Homo sapiens hypothetical protein FLJ10374 (FLJ10374), mRNA
NM 018057	Homo sapiens homolog of rat orphan transporter v7-3 (NTT73), mRNA
NM 018049	Homo sapiens hypothetical protein FLJ10297 (FLJ10297); mRNA
NM 018028	Homo sapiens hypothetical protein FLJ10211 (FLJ10211), mRNA
NM 018000	Homo sapiens hypothetical protein FLJ10116 (FLJ10116), mRNA
NM 016510	Homo sapiens putative selenocysteine lyase (SCLY), mRNA
NM_016434	Homo sapiens tumor necrosis factor receptor superfamily, member 6b, decoy
	(TNFRSF6B), transcript variant 2, mRNA
NM_016289	Homo sapiens MO25 protein (LOC51719), mRNA
NM 016264	Homo sapiens GIOT-2 for gonadotropin inducible transcription repressor-2
	(GIOT-2), mRNA
NM_016149	Homo sapiens protein inhibitor of activated STAT protein PIASy (PIASY), mRNA
NM_015897	Homo sapiens protein inhibitor of activated STAT protein PIASy (PIASY), mRNA
NM 016581	Homo sapiens ECSIT (LOC51295), mRNA
NM 016479	Homo sapiens hypothetical protein (LOC51246), mRNA
NM 016474	Homo sapiens hypothetical protein (LOC51244), mRNA
NM_016094	Homo sapiens HSPC042 protein (LOC51122), mRNA
NM_015942	Homo sapiens CGI-12 protein (LOC51001), mRNA
NM_016475	Homo sapiens hypothetical protein (HSPC213), mRNA
NM_016457	Homo sapiens protein kinase D2 (PKD2), mRNA
NM_016111	Homo sapiens KIAA0683 gene product (KIAA0683), mRNA
NM_014049	Homo sapiens NPD002 protein (NPD002), mRNA
NM_014963	Homo sapiens KIAA0963 protein (KIAA0963), mRNA
NM 015571	Homo sapiens SUMO-1-specific protease (SUSP1), mRNA
NM_014789	Homo sapiens KIAA0628 gene product (KIAA0628), mRNA
NM 014714	Homo sapiens KIAA0590 gene product (KIAA0590), mRNA
NM_014758	Homo sapiens KIAA0254 gene product (KIAA0254), mRNA
NM_014065	Homo sapiens HT001 protein (HT001), mRNA
NM_014170	Homo sapiens HSPC135 protein (HSPC135), mRNA
NM_015462	Homo sapiens DKFZP586L0724 protein (DKFZP586L0724), mRNA
NM_015642	Homo sapiens zinc finger protein 288 (ZNF288), mRNA
NM_015493	Homo sapiens DKFZP434N161 protein (DKFZP434N161), mRNA
NM_014446	Homo sapiens muscle-specific beta 1 integrin binding protein (MIBP), mRNA
NM_013314	Homo sapiens B-cell linker (BLNK), mRNA

NM 007086 Homo sapiens similar to S. pombe dimit + (DiM1), mRNA		
NM 006407 Homo sapiens zinc finger protein 230 (ZNP230), mRNA NM 006087 Homo sapiens RAS-related on chromosome 22 (RRP22), mRNA NM 006087 Homo sapiens grotein-coupled receptor 66 (GPR66), mRNA NM 005815 Homo sapiens Gryotein-coupled receptor 66 (GPR66), mRNA NM 005817 Homo sapiens cargo selection protein (mannose 6 phosphate receptor binding protein) (TIP47), mRNA NM 005818 Homo sapiens protein translation initiation factor (SUI1), mRNA NM 005819 Homo sapiens putative translation initiation factor (SUI1), mRNA NM 005837 Homo sapiens protein (CPR12), mRNA NM 00576 Homo sapiens cornichon-like (CNIL), mRNA NM 004970 Homo sapiens sinutin-like growth factor binding protein, acid labile subunit (IGFALS), mRNA NM 004945 Homo sapiens dynamin 2 (DNM2), mRNA NM 004948 Homo sapiens MADH dehydrogenase (ubiquinone) 1 beta subcomplex, 10 (2kD, PDSW) (NDUFB10), mRNA NM 004124 Homo sapiens glia maturation factor, beta (GMFB), mRNA NM 004124 Homo sapiens glia maturation factor, pamma (GMFG), mRNA NM 004907 Homo sapiens glia maturation factor, pamma (GMFG), mRNA NM 004907 Homo sapiens glia maturation factor, pamma (GMFG), mRNA NM 004917 Homo sapiens protein (ETR101), mRNA NM 004916 Homo sapiens sylaminate early protein (ETR101), mRNA NM 004315 Homo sapiens SP2 transcription factor (SP2), mRNA NM 003406 Homo sapiens syntaxin 10 (STX10), mRNA NM 003110 Homo sapiens spina syntaxin 10 (STX10), mRNA NM 003111 Homo sapiens spina syntaxin 10 (STX10), mRNA NM 00310 Homo sapiens spina syntaxin 10 (STX10), mRNA NM 00311 Homo sapiens spina syntaxin 10 (STX10), mRNA NM 003010 Homo sapiens spina syntaxin 10 (STX10), mRNA NM 003010 Homo sapiens spina syntaxin 10 (STX10), mRNA NM 003010 Homo sapiens spina syntaxin 10 (STX10), mRNA NM 003010 Homo sapiens spina syntaxin 10 (STX10), mRNA NM 003010 Homo sapiens spina syntaxin 10 (STX10), mRNA NM 003010 Homo sapiens spina syntaxin 10 (STX10), mRNA NM 003010 Homo sapiens spina	NM_007086	Homo sapiens AND-1 protein (AND-1), mRNA
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NM_004124	NM_004945	Homo sapiens dynamin 2 (DNM2), mRNA
NM_004124		Homo sapiens RAB3D, member RAS oncogene family (RAB3D), mRNA
NM 004124 Homo sapiens glia maturation factor, beta (GMFB), mRNA NM 004877 Homo sapiens glia maturation factor, gamma (GMFG), mRNA NM 004907 Homo sapiens immediate early protein (ETR101), mRNA NM 004044 Homo sapiens 5-aminorimidazole-4-carboxamide ribonucleotide formyltransferase/IMP cyclohydrolase (ATIC), mRNA NM 004315 Homo sapiens N-acylsphingosine amidohydrolase (acid ceramidase) (ASAH), mRNA NM 004846 Homo sapiens eukaryotic translation initiation factor 4E-like 3 (EIF4EL3), mRNA NM 003765 Homo sapiens syntaxin 10 (STX10), mRNA NM 003110 Homo sapiens Sp2 transcription factor (SP2), mRNA NM 003111 Homo sapiens suclear antigen Sp100 (SP100), mRNA NM 003113 Homo sapiens suclear antigen Sp100 (SP100), mRNA NM 003072 Homo sapiens SWI/SNF related, matrix associated, actin dependent regulator of chromatin, subfamily a, member 4 (SMARCA4), mRNA NM 003072 Homo sapiens proteasome (prosome, macropain) 26S subunit, non-ATPase, 1 (PSMD1), mRNA NM 002704 Homo sapiens pro-platelet basic protein (includes platelet basic protein, beta-thromboglobulin, connective tissue-activating peptide-2) (PPBP), mRNA NM 000889 Homo sapiens attractivating peptide-2) (PPBP), mRNA NM 001687 Homo sapiens BTP synthase, H+ transporting, mitochondrial F1 complex, delta subunit (ATP5D), mRNA NM 032657 Homo sapiens BGF-like module-containing mucin-like receptor EMR3 (EMR3), mRNA NM 032571 Homo sapiens normal mucosa of esophagus specific 1 (NMES1), mRNA NM 032413 Homo sapiens normal mucosa of esophagus specific 1 (NMES1), mRNA NM 015093 Homo sapiens normal mucosa of esophagus specific 1 (NMES1), mRNA NM 031947 Homo sapiens stathmin 1/oncoprotein 18 (STMN1), mRNA	NM_004548	Homo sapiens NADH dehydrogenase (ubiquinone) 1 beta subcomplex, 10 (22kD, PDSW) (NDUFB10), mRNA
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NM_024662 Homo sapiens hypothetical protein FLJ10774 (FLJ10774), mRNA		
		Homo sapiens stathmin 1/oncoprotein 18 (STMN1), mRNA

NM_024637	Homo sapiens beta-galactose-3-O-sulfotransferase, 4 (GAL3ST-4), mRNA
NM_024617	Homo sapiens hypothetical protein FLJ13409 (FLJ13409), mRNA
NM_020796	Homo sapiens sema domain, transmembrane domain (TM), and cytoplasmic domain, (semaphorin) 6A (SEMA6A), mRNA
NM 013283	Homo sapiens methionine adenosyltransferase II, beta (MAT2B), mRNA
	Homo sapiens PR domain containing 2, with ZNF domain (PRDM2), mRNA
NM_012231	
NM_020428	Homo sapiens CTL2 gene (CTL2), mRNA
NM_015866	Homo sapiens PR domain containing 2, with ZNF domain (PRDM2), mRNA
NM_014771	Homo sapiens 95 kDa retinoblastoma protein binding protein; KIAA0661 gene pro (KIAA0661), mRNA
NM 014454	Homo sapiens p53 regulated PA26 nuclear protein (PA26), mRNA
NM_013447	Homo sapiens egf-like module containing, mucin-like, hormone receptor-like sequence 2 (EMR2), mRNA
NM_006499	Homo sapiens lectin, galactoside-binding, soluble, 8 (galectin 8) (LGALS8), mRNA
NM 006031	Homo sapiens pericentrin 2 (kendrin) (PCNT2), mRNA
NM 022040	Homo sapiens Williams-Beuren syndrome chromosome region 5 (WBSCR5),
	transcript variant 1, mRNA
NM_032464	Homo sapiens Williams-Beuren syndrome chromosome region 5 (WBSCR5), transcript variant 4, mRNA
NM_032463	Homo sapiens Williams-Beuren syndrome chromosome region 5 (WBSCR5),
	transcript variant 2, mRNA
NM_014146	Homo sapiens Williams-Beuren syndrome chromosome region 5 (WBSCR5), transcript variant 3, mRNA
NM_031992	Homo sapiens Williams-Beuren syndrome chromosome region 1 (WBSCR1),
1111_031352	transcript variant 2, mRNA
NM_006234	Homo sapiens polymerase (RNA) II (DNA directed) polypeptide J (13.3kD) (POLR2J), transcript variant a, mRNA
NM_032959	Homo sapiens polymerase (RNA) II (DNA directed) polypeptide J (13.3kD) (POLR2J), transcript variant b, mRNA
NM_032958	Homo sapiens polymerase (RNA) II (DNA directed) polypeptide J (13.3kD)
1111_002>00	(POLR2J), transcript variant c, mRNA
NM_002694	Homo sapiens polymerase (RNA) II (DNA directed) polypeptide C (33kD)
	(POLR2C), transcript variant alpha, mRNA
NM 032940	Homo sapiens polymerase (RNA) II (DNA directed) polypeptide C (33kD)
	(POLR2C), transcript variant gamma, mRNA
NM 033011	Homo sapiens plasminogen activator, tissue (PLAT), transcript variant 3, mRNA
NM 000931	Homo sapiens plasminogen activator, tissue (PLAT), transcript variant 2, mRNA
NM 000930	Homo sapiens plasminogen activator, tissue (PLAT), transcript variant 1, mRNA
NM_033013	Homo sapiens nuclear receptor subfamily 1, group I, member 2 (NR1I2),
	transcript variant 3, mRNA
NM 003889	Homo sapiens nuclear receptor subfamily 1, group I, member 2 (NR1I2),
	transcript variant 1, mRNA
NM_022002	Homo sapiens nuclear receptor subfamily 1, group I, member 2 (NR1I2),
	transcript variant 2, mRNA
NM 022170	Homo sapiens Williams-Beuren syndrome chromosome region 1 (WBSCR1),
	transcript variant 1, mRNA
NM 032408.	Homo sapiens bromodomain adjacent to zinc finger domain, 1B (BAZ1B),
	transcript variant 2, mRNA
NM 023005	Homo sapiens bromodomain adjacent to zinc finger domain, 1B (BAZ1B),
	transcript variant 1, mRNA
NM_001024	Homo sapiens ribosomal protein S21 (RPS21), mRNA

NM_012138	Homo sapiens apoptosis antagonizing transcription factor (DED), mRNA
NM_016343	Homo sapiens centromere protein F (350/400kD, mitosin) (CENPF), mRNA
NM_032988	Homo sapiens transducin (beta)-like 2 (TBL2), transcript variant 2, mRNA
NM_032052	Homo sapiens zinc finger protein 278 (ZNF278), transcript variant 3, mRNA
NM_032051	Homo sapiens zinc finger protein 278 (ZNF278), transcript variant 4, mRNA
NM_032050	Homo sapiens zinc finger protein 278 (ZNF278), transcript variant 2, mRNA
NM_014323	Homo sapiens zinc finger protein 278 (ZNF278), transcript variant 1, mRNA
NM_033003	Homo sapiens general transcription factor II, i (GTF2I), transcript variant 5,
	mRNA
NM_001518	Homo sapiens general transcription factor II, i (GTF2I), transcript variant 4,
	mRNA
NM_033001	Homo sapiens general transcription factor II, i (GTF2I), transcript variant 3,
	mRNA .
NM_033000	Homo sapiens general transcription factor II, i (GTF2I), transcript variant 2, mRNA
NM_032999	Homo sapiens general transcription factor II, i (GTF2I), transcript variant 1, mRNA
NM_002904	Homo sapiens RD RNA-binding protein (RDBP), mRNA
NM_002755	Homo sapiens mitogen-activated protein kinase kinase 1 (MAP2K1), mRNA
NM_012453	Homo sapiens transducin (beta)-like 2 (TBL2), transcript variant 1, mRNA
NM_006347	Homo sapiens peptidyl prolyl isomerase H (cyclophilin H) (PPIH), mRNA
NM_001631	Homo sapiens alkaline phosphatase, intestinal (ALPI), mRNA
NM_021151	Homo sapiens carnitine O-octanoyltransferase (CROT), mRNA
NM_005090	Homo sapiens phospholipase A2, group IVB (cytosolic) (PLA2G4B), mRNA
NM_000124	Homo sapiens excision repair cross-complementing rodent repair deficiency,
	complementation group 6 (ERCC6), mRNA
NM_020157	Homo sapiens otoraplin (OTOR), mRNA
NM_018313	Homo sapiens polybromo 1 (PB1), mRNA
NM_018165	Homo sapiens polybromo 1 (PB1), mRNA
NM_016503	Homo sapiens mitochondrial ribosomal protein L30 (MRPL30), mRNA
NM_012139	Homo sapiens deafness locus associated putative guanine nucleotide exchange f
	(DELGEF), mRNA
NM_007061	Homo sapiens serum constituent protein (MSE55), mRNA
NM_005379	Homo sapiens myosin IA (MYO1A), mRNA
NM_000500	Homo sapiens cytochrome P450, subfamily XXIA (steroid 21-hydroxylase,
37.5 00000	congenital adrenal hyperplasia), polypeptide 2 (CYP21A2), mRNA
NM_000063	Homo sapiens complement component 2 (C2), mRNA
NM_014078	Homo sapiens mitochondrial ribosomal protein L13 (MRPL13), mRNA
NM_021134	Homo sapiens mitochondrial ribosomal protein L23 (MRPL23), mRNA
NM_020249	Homo sapiens a disintegrin-like and metalloprotease (reprolysin type) with
3D 6 010006	thrombospondin type 1 motif, 9 (ADAMTS9), mRNA
NM_018094	Homo sapiens G1 to S phase transition 2 (GSPT2), mRNA
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NM_014180	Homo sapiens mitochondrial ribosomal protein L22 (MRPL22), mRNA
NM_014175	Homo sapiens mitochondrial ribosomal protein L15 (MRPL15), mRNA
NM_014175 NM_015385	Homo sapiens mitochondrial ribosomal protein L15 (MRPL15), mRNA Homo sapiens SH3-domain protein 5 (ponsin) (SH3D5), mRNA
NM_014175 NM_015385 NM_006434	Homo sapiens mitochondrial ribosomal protein L15 (MRPL15), mRNA Homo sapiens SH3-domain protein 5 (ponsin) (SH3D5), mRNA Homo sapiens SH3-domain protein 5 (ponsin) (SH3D5), mRNA
NM_014175 NM_015385 NM_006434 NM_000135	Homo sapiens mitochondrial ribosomal protein L15 (MRPL15), mRNA Homo sapiens SH3-domain protein 5 (ponsin) (SH3D5), mRNA Homo sapiens SH3-domain protein 5 (ponsin) (SH3D5), mRNA Homo sapiens Fanconi anemia, complementation group A (FANCA), mRNA
NM 014175 NM 015385 NM 006434 NM 000135 NM 005656	Homo sapiens mitochondrial ribosomal protein L15 (MRPL15), mRNA Homo sapiens SH3-domain protein 5 (ponsin) (SH3D5), mRNA Homo sapiens SH3-domain protein 5 (ponsin) (SH3D5), mRNA Homo sapiens Fanconi anemia, complementation group A (FANCA), mRNA Homo sapiens transmembrane protease, serine 2 (TMPRSS2), mRNA
NM 014175 NM 015385 NM 006434 NM 000135 NM 005656 NM 021974	Homo sapiens mitochondrial ribosomal protein L15 (MRPL15), mRNA Homo sapiens SH3-domain protein 5 (ponsin) (SH3D5), mRNA Homo sapiens SH3-domain protein 5 (ponsin) (SH3D5), mRNA Homo sapiens Fanconi anemia, complementation group A (FANCA), mRNA Homo sapiens transmembrane protease, serine 2 (TMPRSS2), mRNA Homo sapiens polymerase (RNA) II (DNA directed) polypeptide F (POLR2F), mRNA
NM 014175 NM 015385 NM 006434 NM 000135 NM 005656	Homo sapiens mitochondrial ribosomal protein L15 (MRPL15), mRNA Homo sapiens SH3-domain protein 5 (ponsin) (SH3D5), mRNA Homo sapiens SH3-domain protein 5 (ponsin) (SH3D5), mRNA Homo sapiens Fanconi anemia, complementation group A (FANCA), mRNA Homo sapiens transmembrane protease, serine 2 (TMPRSS2), mRNA Homo sapiens polymerase (RNA) II (DNA directed) polypeptide F (POLR2F),

NM 032965 Homo sapiens small inducible cytokine subfamily A (Cys-Cys), member 15 (SCYA15), transcript variant 1, mRNA		
NM_032964 Homo sapiens small inducible cytokine subfamily A (Cys-Cys), member 15 (SCYA15), transcript variant 1, mRNA NM_032454 Homo sapiens serine/threonine kinase 19 (STK19), transcript variant 2, mRNA NM_03297 Homo sapiens ZW10 interactor (ZWINT), transcript variant 1, mRNA NM_03297 Homo sapiens translocation protein 1 (TLOCI), mRNA NM_03247 Homo sapiens translocation protein 1 (TLOCI), mRNA NM_03247 Homo sapiens translocation protein 1 (TLOCI), mRNA NM_03247 Homo sapiens small inducible cytokine subfamily A (Cys-Cys), member 14 (SCYA14), transcript variant 1, mRNA NM_032961 Homo sapiens small inducible cytokine subfamily A (Cys-Cys), member 14 (SCYA14), transcript variant 3, mRNA NM_032962 Homo sapiens small inducible cytokine subfamily A (Cys-Cys), member 14 (SCYA14), transcript variant 2, mRNA NM_032962 Homo sapiens small inducible cytokine subfamily A (Cys-Cys), member 14 (SCYA14), transcript variant 2, mRNA Homo sapiens programmed cell death 4 (neoplastic transformation inhibitor) (PDCD4), mRNA Homo sapiens programmed cell death 4 (neoplastic transformation inhibitor) (PDCD4), mRNA Homo sapiens SEC63, endoplasmic reticulum translocon component (S. cerevisiae (SEC631L), mRNA NM_001028 Homo sapiens protein translocation complex beta (SEC61B), mRNA NM_001028 Homo sapiens ribosomal protein S25 (RPS25), mRNA Homo sapiens ribosomal protein S19 (RPS19), mRNA Homo sapiens ribosomal protein S16 (RPS16), mRNA Homo sapiens ribosomal protein S16 (RPS16), mRNA Homo sapiens ribosomal protein S16 (RPS16), mRNA Homo sapiens ribosomal protein S17 (RPS17), mRNA Homo sapiens RecQ protein-like (DNA helicase Q1-like) (RECQL), transcript variant 2, mRNA Homo sapiens polymerase (RNA) II (DNA directed) polypeptide L (7.6kD) (POLR21), mRNA Homo sapiens polymerase (RNA) II (DNA directed) polypeptide D (POLR2D)	NM_032965	Homo sapiens small inducible cytokine subfamily A (Cys-Cys), member 15 (SCYA15), transcript variant 3, mRNA
(SCYA15), transcript variant 1, mRNA NM 032454 Homo sapiens serine/threonine kinase 19 (STK19), transcript variant 2, mRNA NM 032997 Homo sapiens ZW10 interactor (ZWINT), transcript variant 1, mRNA NM 032907 Homo sapiens translocation protein 1 (TLOC1), mRNA NM 032470 Homo sapiens translocation protein 1 (TLOC1), mRNA NM 032470 Homo sapiens transcrin XB (TNXB), transcript variant XB-S, mRNA NM_04166 Homo sapiens small inducible cytokine subfamily A (Cys-Cys), member 14 (SCYA14), transcript variant 1, mRNA NM_032963 Homo sapiens small inducible cytokine subfamily A (Cys-Cys), member 14 (SCYA14), transcript variant 3, mRNA NM_032962 Homo sapiens small inducible cytokine subfamily A (Cys-Cys), member 14 (SCYA14), transcript variant 3, mRNA NM_041219 Homo sapiens small inducible cytokine subfamily A (Cys-Cys), member 14 (SCYA14), transcript variant 2, mRNA NM_041219 Homo sapiens small inducible cytokine subfamily A (Cys-Cys), member 14 (SCYA14), transcript variant 2, mRNA NM_004197 Homo sapiens protein translocation of transformation inhibitor (PDCD4), mRNA NM_004197 Homo sapiens sprinder threonine kinase 19 (STK19), transcript variant 1, mRNA NM_001021 Homo sapiens sprinder threonine kinase 19 (STK19), transcript variant 1, mRNA NM_001028 Homo sapiens protein translocation complex beta (SEC61B), mRNA NM_001029 Homo sapiens ribosomal protein S19 (RPS19), mRNA NM_001020 Homo sapiens ribosomal protein S19 (RPS19), mRNA NM_001021 Homo sapiens ribosomal protein S16 (RPS16), mRNA NM_001021 Homo sapiens ribosomal protein S16 (RPS16), mRNA NM_001023 Homo sapiens ribosomal protein S18 (RPS13), mRNA NM_001243 Homo sapiens ribosomal protein S18 (RPS13), mRNA NM_002907 Homo sapiens ribosomal protein S18 (RPS13), mRNA NM_002907 Homo sapiens secQ protein-like (DNA helicase Q1-like) (RECQL), transcript variant 1, mRNA NM_002907 Homo sapiens polymerase (RNA) II (DNA directed) polypeptide L (7.6kD) (POLR2D), mRNA NM_002903 Homo sapiens polymerase (RNA) II (DNA directed) polypeptide D	NM 032964	
NM 032971 Homo sapiens ZW10 interactor (ZWINT), transcript variant 1, mRNA NM 032972 Homo sapiens ZW10 interactor (ZWINT), transcript variant 2, mRNA NM 032402 Homo sapiens translocation protein 1 (TLOC1), mRNA NM 032410 Homo sapiens transcrin XB (TNXB), transcript variant XB-S, mRNA NM 032410 Homo sapiens transcrin XB (TNXB), transcript variant XB-S, mRNA NM 032410 Homo sapiens small inducible cytokine subfamily A (Cys-Cys), member 14 (SCYA14), transcript variant 1, mRNA NM 032963 Homo sapiens small inducible cytokine subfamily A (Cys-Cys), member 14 (SCYA14), transcript variant 2, mRNA NM 032962 Homo sapiens small inducible cytokine subfamily A (Cys-Cys), member 14 (SCYA14), transcript variant 2, mRNA NM 021219 Homo sapiens simall inducible cytokine subfamily A (Cys-Cys), member 14 (SCYA14), transcript variant 2, mRNA NM 021219 Homo sapiens programmed cell death 4 (neoplastic transformation inhibitor) (PDCD4), mRNA NM 001419 Homo sapiens programmed cell death 4 (neoplastic transformation inhibitor) (PDCD4), mRNA NM 0004197 Homo sapiens scine/threonine kinase 19 (STK19), transcript variant 1, mRNA NM 007214 Homo sapiens scine/threonine kinase 19 (STK19), transcript variant 1, mRNA NM 001028 Homo sapiens protein Eranslocation complex beta (SEC61B), mRNA NM 001028 Homo sapiens ribosomal protein S25 (RPS25), mRNA NM 001020 Homo sapiens ribosomal protein S19 (RPS19), mRNA NM 001021 Homo sapiens ribosomal protein S16 (RPS19), mRNA NM 001021 Homo sapiens ribosomal protein S16 (RPS16), mRNA NM 001017 Homo sapiens ribosomal protein S18 (RPS13), mRNA NM 001024 Homo sapiens ribosomal protein S18 (RPS13), mRNA NM 001024 Homo sapiens RecQ protein-like (DNA helicase Q1-like) (RECQL), transcript variant 1, mRNA NM 002907 Homo sapiens polymerase (RNA) II (DNA directed) polypeptide L (7.6kD) (POLR2L), mRNA NM 00623 Homo sapiens polymerase (RNA) II (DNA directed) polypeptide D (POLR2D), mRNA NM 002695 Homo sapiens polymerase (RNA) II (DNA directed) polypeptide A (220kD) (POLR2L), mRNA NM 0032971 Homo sapiens protocadherin 22 (
NM_032297 Homo sapiens ZW10 interactor (ZWINT), transcript variant 2, mRNA NM_003261 Homo sapiens translocation protein I (TLOC1), mRNA NM_032470 Homo sapiens translocation protein I (TLOC1), mRNA NM_032470 Homo sapiens small inducible cytokine subfamily A (Cys-Cys), member 14 (SCYA14), transcript variant 1, mRNA NM_032963 Homo sapiens small inducible cytokine subfamily A (Cys-Cys), member 14 (SCYA14), transcript variant 3, mRNA NM_032962 Homo sapiens small inducible cytokine subfamily A (Cys-Cys), member 14 (SCYA14), transcript variant 3, mRNA NM_032962 Homo sapiens small inducible cytokine subfamily A (Cys-Cys), member 14 (SCYA14), transcript variant 2, mRNA NM_01419 Homo sapiens junctional adhesion molecule 2 (JAM2), mRNA NM_014456 Homo sapiens programmed cell death 4 (neoplastic transformation inhibitor) (PDCD4), mRNA NM_004197 Homo sapiens serine/threonine kinase 19 (STK19), transcript variant 1, mRNA NM_007214 Homo sapiens SEC63, endoplasmic reticulum translocon component (S. cerevisiae (SEC631), mRNA NM_001028 Homo sapiens ribosomal protein S25 (RPS25), mRNA NM_001028 Homo sapiens ribosomal protein S19 (RPS19), mRNA NM_001020 Homo sapiens ribosomal protein S19 (RPS19), mRNA NM_001020 Homo sapiens ribosomal protein S19 (RPS19), mRNA NM_001021 Homo sapiens ribosomal protein S16 (RPS16), mRNA NM_001021 Homo sapiens ribosomal protein S16 (RPS16), mRNA NM_001021 Homo sapiens ribosomal protein S18 (RPS17), mRNA NM_001021 Homo sapiens ribosomal protein S18 (RPS13), mRNA NM_001021 Homo sapiens ribosomal protein S18 (RPS13), mRNA NM_001023 Homo sapiens RecQ protein-like (DNA helicase Q1-like) (RECQL), transcript variant 1, mRNA NM_002907 Homo sapiens RecQ protein-like (DNA helicase Q1-like) (RECQL), transcript variant 1, mRNA NM_002907 Homo sapiens polymerase (RNA) II (DNA directed) polypeptide H (POLR2H), mRNA NM_002907 Homo sapiens polymerase (RNA) II (DNA directed) polypeptide D (POLR2D), mRNA NM_002907 Homo sapiens polymerase (RNA) II (DNA directed) polypeptide D (POLR2D), mRNA NM_002907 Homo sapiens protocad	NM_032454	Homo sapiens serine/threonine kinase 19 (STK19), transcript variant 2, mRNA
NM 032402 Homo sapiens translocation protein 1 (TLOC1), mRNA NM 032470 Homo sapiens small inducible cytokine subfamily A (Cys-Cys), member 14 (SCYA14), transcript variant 1, mRNA NM_032963 Homo sapiens small inducible cytokine subfamily A (Cys-Cys), member 14 (SCYA14), transcript variant 3, mRNA NM_032962 Homo sapiens small inducible cytokine subfamily A (Cys-Cys), member 14 (SCYA14), transcript variant 3, mRNA NM_032962 Homo sapiens small inducible cytokine subfamily A (Cys-Cys), member 14 (SCYA14), transcript variant 3, mRNA NM_032962 Homo sapiens small inducible cytokine subfamily A (Cys-Cys), member 14 (SCYA14), transcript variant 2, mRNA NM_032962 Homo sapiens small inducible cytokine subfamily A (Cys-Cys), member 14 (SCYA14), transcript variant 2, mRNA NM_014456 Homo sapiens programmed cell death 4 (neoplastic transformation inhibitor) (PDCD4), mRNA NM_001497 Homo sapiens srine/threonine kinase 19 (STK19), transcript variant 1, mRNA NM_007214 Homo sapiens sribectine translocation complex beta (SEC61B), mRNA NM_001028 Homo sapiens protein translocation complex beta (SEC61B), mRNA NM_001020 Homo sapiens ribosomal protein S15 (RPS19), mRNA NM_001021 Homo sapiens ribosomal protein S16 (RPS16), mRNA NM_001021 Homo sapiens ribosomal protein S16 (RPS16), mRNA NM_001010 Homo sapiens ribosomal protein S16 (RPS16), mRNA NM_001017 Homo sapiens ribosomal protein S16 (RPS16), mRNA NM_001017 Homo sapiens ribosomal protein S16 (RPS16), mRNA NM_001018 Homo sapiens ribosomal protein S16 (RPS16), mRNA NM_002907 Homo sapiens ribosomal protein S16 (RPS16), mRNA NM_002901 Homo sapiens ribosomal protein S16 (RPS16), mRNA NM_002901 Homo sapiens RecQ protein-like (DNA helicase Q1-like) (RECQL), transcript variant 2, mRNA NM_002901 Homo sapiens RecQ protein-like (DNA helicase Q1-like) (RECQL), transcript variant 2, mRNA NM_002901 Homo sapiens polymerase (RNA) II (DNA directed) polypeptide L (7.6kD) (POLR2L), mRNA NM_002902 Homo sapiens polymerase (RNA) II (DNA directed) polypeptide D (POLR2D), mRNA NM_002903 Homo sapiens protocad	NM_007057	Homo sapiens ZW10 interactor (ZWINT), transcript variant 1, mRNA
NM 032470 Homo sapiens tenascin XB (TNXB), transcript variant XB-S, mRNA NM 004166 Homo sapiens small inducible cytokine subfamily A (Cys-Cys), member 14 (SCYA14), transcript variant 1, mRNA NM 032963 Homo sapiens small inducible cytokine subfamily A (Cys-Cys), member 14 (SCYA14), transcript variant 3, mRNA NM 032962 Homo sapiens small inducible cytokine subfamily A (Cys-Cys), member 14 (SCYA14), transcript variant 2, mRNA NM 021219 Homo sapiens protein transladesion molecule 2 (JAM2), mRNA NM 014185 Homo sapiens protein transladesion molecule 2 (JAM2), mRNA NM 004197 Homo sapiens serine/threonine kinase 19 (STK19), transcript variant 1, mRNA NM 004197 Homo sapiens serine/threonine kinase 19 (STK19), transcript variant 1, mRNA NM 004197 Homo sapiens serine/threonine kinase 19 (STK19), transcript variant 1, mRNA NM 001028 Homo sapiens protein translocation complex beta (SEC61B), mRNA NM 001028 Homo sapiens protein translocation complex beta (SEC61B), mRNA NM 001028 Homo sapiens ribosomal protein S19 (RPS19), mRNA NM 001020 Homo sapiens ribosomal protein S19 (RPS19), mRNA NM 001010 Homo sapiens ribosomal protein S16 (RPS16), mRNA NM 001018 Homo sapiens ribosomal protein S16 (RPS16), mRNA NM 001012 Homo sapiens ribosomal protein S16 (RPS16), mRNA NM 001012 Homo sapiens ribosomal protein S18 (RPS13), mRNA NM 001012 Homo sapiens ribosomal protein S18 (RPS13), mRNA NM 001017 Homo sapiens RecQ protein-like (DNA helicase Q1-like) (RECQL), transcript variant 1, mRNA NM 002907 Homo sapiens RecQ protein-like (DNA helicase Q1-like) (RECQL), transcript variant 2, mRNA NM 0021128 Homo sapiens polymerase (RNA) II (DNA directed) polypeptide L (7.6kD) (POLR2L), mRNA NM 006232 Homo sapiens polymerase (RNA) II (DNA directed) polypeptide E (25kD) (POLR2L), mRNA NM 004805 Homo sapiens polymerase (RNA) II (DNA directed) polypeptide E (25kD) (POLR2L), mRNA NM 004805 Homo sapiens polymerase (RNA) II (DNA directed) polypeptide E (25kD) (POLR2L), mRNA NM 003937 Homo sapiens protocadherin 22 (PCDH22), transcript variant c, mRNA NM 003971	NM_032997	Homo sapiens ZW10 interactor (ZWINT), transcript variant 2, mRNA
NM_004166 Homo sapiens small inducible cytokine subfamily A (Cys-Cys), member 14 (SCYA14), transcript variant 1, mRNA (SCYA14), transcript variant 3, mRNA (Cys-Cys), member 14 (SCYA14), transcript variant 3, mRNA (Cys-Cys), member 14 (SCYA14), transcript variant 3, mRNA (Cys-Cys), member 14 (SCYA14), transcript variant 2, mRNA (SCYA14), transcript variant 3, mRNA (Cys-Cys), member 14 (SCYA14), transcript variant 2, mRNA (SCYA14), transcript variant 1, mRNA (SCYA14), transcript variant 2, mRNA (SCYA14), transcript variant 3, mRNA (SCYA144), transcript variant 3, mRNA (SCYA144), transcript variant 3, mRNA (SCYA144), transcript variant 4, mRNA (SCYA144), transcript variant 5, mRNA (SCYA144), transcript variant 6, mRNA (SCYA144), transcr	NM_003262	Homo sapiens translocation protein 1 (TLOC1), mRNA
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		Homo sapiens protocadherin 9 (PCDH9), mRNA
NM_032949 Homo sapiens protocadherin 8 (PCDH8), transcript variant 2, mRNA		
	NM_032949	Homo sapiens protocadherin 8 (PCDH8), transcript variant 2, mRNA

NM_032457	Homo sapiens BH-protocadherin (brain-heart) (PCDH7), transcript variant c, mRNA
NM_032456	Homo sapiens BH-protocadherin (brain-heart) (PCDH7), transcript variant b, mRNA
NM_002589	Homo sapiens BH-protocadherin (brain-heart) (PCDH7), transcript variant a, mRNA
NM 016580	Homo sapiens protocadherin 12 (PCDH12), mRNA
NM 032420	Homo sapiens protocadherin 1 (cadherin-like 1) (PCDH1), transcript variant 2,
	mRNA
NM_032969	Homo sapiens protocadherin 11 (PCDH11), transcript variant d, mRNA
NM_032968	Homo sapiens protocadherin 11 (PCDH11), transcript variant c. mRNA
NM_032967	Homo sapiens protocadherin 11 (PCDH11), transcript variant b, mRNA
NM_032950	Homo sapiens matrix metalloproteinase 28 (MMP28), transcript variant 2, mRNA
NM_024302	Homo sapiens matrix metalloproteinase 28 (MMP28), transcript variant 1, mRNA
NM_006575	Homo sapiens mitogen-activated protein kinase kinase kinase kinase 5
	(MAP4K5), mRNA
NM_004635	Homo sapiens mitogen-activated protein kinase-activated protein kinase 3 (MAPKAPK3), mRNA
NM_002587	Homo sapiens protocadherin 1 (cadherin-like 1) (PCDH1), transcript variant 1, mRNA
NM 004759	
14141_004739	Homo sapiens mitogen-activated protein kinase-activated protein kinase 2 (MAPKAPK2), transcript variant 1, mRNA
NM_032960	Homo sapiens mitogen-activated protein kinase-activated protein kinase 2
	(MAPKAPK2), transcript variant 2, mRNA
NM_032515	Homo sapiens Bcl-2-related ovarian killer protein-like (BOKL), mRNA
NM_015166	Homo sapiens KIAA0027 protein (MLC1), mRNA
NM_001795	Homo sapiens cadherin 5, type 2, VE-cadherin (vascular epithelium) (CDH5), mRNA
NM_001794	Homo sapiens cadherin 4, type 1, R-cadherin (retinal) (CDH4), mRNA
NM 001793	Homo sapiens cadherin 3, type 1, P-cadherin (placental) (CDH3), mRNA
NM 001792	Homo sapiens cadherin 2, type 1, N-cadherin (neuronal) (CDH2), mRNA
NM 004360	Homo sapiens cadherin 1, type 1, E-cadherin (epithelial) (CDH1), mRNA
NM 006137	Homo sapiens CD7 antigen (p41) (CD7), mRNA
NM 005864	Homo sapiens signal transduction protein (SH3 containing) (EFS2), transcript
	variant 1, mRNA
NM_032459	Homo sapiens signal transduction protein (SH3 containing) (EFS2), transcript variant 2, mRNA
NM_032107	Homo sapiens lethal (3) malignant brain tumor l(3)mbt protein (Drosophila) ho
	(H-L(3)MBT), transcript variant II, mRNA
NM_015478	Homo sapiens lethal (3) malignant brain tumor l(3)mbt protein (Drosophila) ho
	(H-L(3)MBT), transcript variant I, mRNA
NM_004318	Homo sapiens aspartate beta-hydroxylase (ASPH), transcript variant 1, mRNA
NM_032468	Homo sapiens aspartate beta-hydroxylase (ASPH), transcript variant 2, mRNA
NM_032467	Homo sapiens aspartate beta-hydroxylase (ASPH), transcript variant 4, mRNA
NM_032466	Homo sapiens aspartate beta-hydroxylase (ASPH), transcript variant 3, mRNA
NM_020164	Homo sapiens aspartate beta-hydroxylase (ASPH), transcript variant 5, mRNA
NM_014217	Homo sapiens potassium channel, subfamily K, member 2 (TREK-1) (KCNK2), mRNA
NM_031498	Homo sapiens guanine nucleotide binding protein (G protein), gamma
	transducing activity polypeptide 2 (GNGT2), mRNA
	p-yp-prido 2 (01.012), million

37.5	
NM_031311	Homo sapiens carboxypeptidase, vitellogenic-like (CPVL), mRNA
NM_022768	Homo sapiens RNA binding motif protein 15 (RBM15), mRNA
NM_021797	Homo sapiens eosinophil chemotactic cytokine (TSA1902), mRNA
NM_014330	Homo sapiens protein phosphatase 1, regulatory (inhibitor) subunit 15A (PPP1R15A), mRNA
NM_014522	Homo sapiens protocadherin 11 (PCDH11), transcript variant a, mRNA
NM_003004	Homo sapiens secreted and transmembrane 1 (SECTM1), mRNA
NM_002696	Homo sapiens polymerase (RNA) II (DNA directed) polypeptide G (POLR2G), mRNA
NM_000938	Homo sapiens polymerase (RNA) II (DNA directed) polypeptide B (140kD) (POLR2B), mRNA
NM_001372	Homo sapiens dynein, axonemal, heavy polypeptide 9 (DNAH9), transcript variant 2, mRNA
NM_004215	Homo sapiens estrogen receptor binding site associated, antigen, 9 (EBAG9), mRNA
NM_005111	Homo sapiens crystallin, zeta (quinone reductase)-like 1 (CRYZL1), mRNA
NM_004381	Homo sapiens cAMP responsive element binding protein-like 1 (CREBL1), mRNA
NM_000592	Homo sapiens complement component 4B (C4B), mRNA
NM_007293	Homo sapiens complement component 4A (C4A), mRNA
NM_032603	Homo sapiens lysyl oxidase-like 3 (LOXL3), mRNA
NM_023937	Homo sapiens mitochondrial ribosomal protein L34 (MRPL34), mRNA
NM_022567	Homo sapiens nyctalopin (NYX), mRNA
NM_022467	Homo sapiens carbohydrate (N-acetylgalactosamine 4-0) sulfotransferase 8 (CHST8), mRNA
NM_016557	Homo sapiens orphan seven-transmembrane receptor, chemokine related (VSHK1), mRNA
NM_016116	Homo sapiens ankyrin repeat and SOCS box-containing 4 (ASB4), mRNA
NM_016114	Homo sapiens ankyrin repeat and SOCS box-containing 1 (ASB1), mRNA
NM_016115	Homo sapiens ankyrin repeat and SOCS box-containing 3 (ASB3), mRNA
NM_014398	Homo sapiens lysosomal-associated membrane protein 3 (LAMP3), mRNA
NM_014434	Homo sapiens NADPH-dependent FMN and FAD containing oxidoreductase (NR1), mRNA
NM_004860	Homo sapiens fragile X mental retardation, autosomal homolog 2 (FXR2), mRNA
NM_006850	Homo sapiens interleukin 24 (IL24), mRNA
NM_006541	Homo sapiens thioredoxin-like 2 (TXNL2), mRNA
NM_004662	Homo sapiens dynein, axonemal, heavy polypeptide 9 (DNAH9), transcript variant 1, mRNA
NM_000029	Homo sapiens angiotensinogen (serine (or cysteine) proteinase inhibitor, clade A (alpha-1 antiproteinase, antitrypsin), member 8) (AGT), mRNA
NM_004050	Homo sapiens BCL2-like 2 (BCL2L2), mRNA
NM_004049	Homo sapiens BCL2-related protein A1 (BCL2A1), mRNA
NM_001623	Homo sapiens allograft inflammatory factor 1 (AIF1), transcript variant 3, mRNA
NM_032955	Homo sapiens allograft inflammatory factor 1 (AIF1), transcript variant 1, mRNA
NG_000010	Homo sapiens genomic cytochrome P450, subfamily IIA (phenobarbital-inducible) (CYP2A.2@) on chromosome 19
NM_004847	Homo sapiens allograft inflammatory factor 1 (AIF1), transcript variant 2, mRNA
NM_005452	Homo sapiens chromosome 6 open reading frame 11 (C6orf11), mRNA
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NM_031282	Homo sapiens immunoglobulin superfamily receptor translocation associated 1 (IRTA1), mRNA
NM_031281	Homo sapiens immunoglobulin superfamily receptor translocation associated 2 (IRTA2), mRNA
NM_000767	Homo sapiens cytochrome P450, subfamily IIB (phenobarbital-inducible), polypeptide 6 (CYP2B6), mRNA
NM 020165	Homo sapiens postreplication repair protein hRAD18p (RAD18), mRNA
NM 001710	Homo sapiens B-factor, properdin (BF), mRNA
NM 021800	Homo sapiens J domain containing protein 1 (JDP1), mRNA
NM 020404	Homo sapiens tumor endothelial marker 1 precursor (TEM1), mRNA
NM_006672	Homo sapiens solute carrier family 22 (organic anion transporter), member 7
14141_000072	(SLC22A7), mRNA
NM 006398	Homo sapiens diubiquitin (UBD), mRNA
NM_005445	Homo sapiens chondroitin sulfate proteoglycan 6 (bamacan) (CSPG6), mRNA
NM 017495	Homo sapiens enontrioun sunate proteogrycan o (banacian) (CSPGO), mRNA
NM 001632	Homo sapiens seed (AISKNASES), inklyA Homo sapiens alkaline phosphatase, placental (Regan isozyme) (ALPP), mRNA
NM 030773	Homo sapiens beta tubulin 1, class VI (TUBB1), mRNA
NM 020643	Homo sapiens chromosome 11 open reading frame 16 (C11orf16), mRNA
NM 020644	Homo sapiens chromosome 11 open reading frame 15 (C11orf15), mRNA Homo sapiens chromosome 11 open reading frame 15 (C11orf15), mRNA
NM 020642	Homo sapiens chromosome 11 open reading frame 17 (C11orf17), mRNA
NM_020201	Homo sapiens 5' nucleotidase, mitochondrial (NT5M), mRNA
NM 003203	Homo sapiens chromosome 2 open reading frame 3 (C2orf3), mRNA
NM 007175	Homo sapiens chromosome 8 open reading frame 2 (C8orf2), mRNA
NM_007023	Homo sapiens cAMP-regulated guanine nucleotide exchange factor II (CAMP-
	GEFII), mRNA
NM_006589	Homo sapiens chromosome 1 open reading frame 2 (Clorf2), mRNA
NM_006105	Homo sapiens Rap1 guanine-nucleotide-exchange factor directly activated by cA (EPAC), mRNA
NM_005637	Homo sapiens synovial sarcoma translocation, chromosome 18 (SS18), mRNA
NM_001213	Homo sapiens chromosome 1 open reading frame 1 (C1orf1), mRNA
NM_002354	Homo sapiens tumor-associated calcium signal transducer 1 (TACSTD1),
NM_003492	mRNA

NM_003797	Homo sapiens chromosome X open reading frame 12 (CXorf12), mRNA
NM_003797 NM_032863	Homo sapiens chromosome X open reading frame 12 (CXorf12), mRNA Homo sapiens embryonic ectoderm development (EED), mRNA
	Homo sapiens chromosome X open reading frame 12 (CXorf12), mRNA Homo sapiens embryonic ectoderm development (EED), mRNA Homo sapiens hypothetical protein FLJ14927 (FLJ14927), mRNA
NM_032863	Homo sapiens chromosome X open reading frame 12 (CXorf12), mRNA Homo sapiens embryonic ectoderm development (EED), mRNA
NM_032863 NM_032813	Homo sapiens chromosome X open reading frame 12 (CXorf12), mRNA Homo sapiens embryonic ectoderm development (EED), mRNA Homo sapiens hypothetical protein FLJ14927 (FLJ14927), mRNA Homo sapiens hypothetical protein FLJ14624 (FLJ14624), mRNA Homo sapiens myopalladin (FLJ14437), mRNA
NM_032863 NM_032813 NM_032578	Homo sapiens chromosome X open reading frame 12 (CXorf12), mRNA Homo sapiens embryonic ectoderm development (EED), mRNA Homo sapiens hypothetical protein FLJ14927 (FLJ14927), mRNA Homo sapiens hypothetical protein FLJ14624 (FLJ14624), mRNA
NM 032863 NM 032813 NM 032578 NM 032385	Homo sapiens chromosome X open reading frame 12 (CXorf12), mRNA Homo sapiens embryonic ectoderm development (EED), mRNA Homo sapiens hypothetical protein FLJ14927 (FLJ14927), mRNA Homo sapiens hypothetical protein FLJ14624 (FLJ14624), mRNA Homo sapiens myopalladin (FLJ14437), mRNA Homo sapiens chromosome 5 open reading frame 4 (C5orf4), mRNA Homo sapiens hypothetical protein FLJ23511 (FLJ23511), mRNA Homo sapiens chromosome 9 open reading frame 5 (C9orf5), mRNA
NM 032863 NM 032813 NM 032578 NM 032385 NM 032239	Homo sapiens chromosome X open reading frame 12 (CXorf12), mRNA Homo sapiens embryonic ectoderm development (EED), mRNA Homo sapiens hypothetical protein FLJ14927 (FLJ14927), mRNA Homo sapiens hypothetical protein FLJ14624 (FLJ14624), mRNA Homo sapiens myopalladin (FLJ14437), mRNA Homo sapiens chromosome 5 open reading frame 4 (C5orf4), mRNA Homo sapiens hypothetical protein FLJ23511 (FLJ23511), mRNA
NM 032863 NM 032813 NM 032578 NM 032385 NM 032239 NM 032012	Homo sapiens chromosome X open reading frame 12 (CXorf12), mRNA Homo sapiens embryonic ectoderm development (EED), mRNA Homo sapiens hypothetical protein FLJ14927 (FLJ14927), mRNA Homo sapiens hypothetical protein FLJ14624 (FLJ14624), mRNA Homo sapiens myopalladin (FLJ14437), mRNA Homo sapiens chromosome 5 open reading frame 4 (C5orf4), mRNA Homo sapiens hypothetical protein FLJ23511 (FLJ23511), mRNA Homo sapiens chromosome 9 open reading frame 5 (C9orf5), mRNA
NM 032863 NM 032813 NM 032578 NM 032385 NM 032239 NM 032012 NM 031922	Homo sapiens chromosome X open reading frame 12 (CXorf12), mRNA Homo sapiens embryonic ectoderm development (EED), mRNA Homo sapiens hypothetical protein FLJ14927 (FLJ14927), mRNA Homo sapiens hypothetical protein FLJ14624 (FLJ14624), mRNA Homo sapiens myopalladin (FLJ14437), mRNA Homo sapiens chromosome 5 open reading frame 4 (C5orf4), mRNA Homo sapiens hypothetical protein FLJ23511 (FLJ23511), mRNA Homo sapiens chromosome 9 open reading frame 5 (C9orf5), mRNA Homo sapiens RALBP1 protein (LOC83859), mRNA Homo sapiens cat eye syndrome chromosome region, candidate 6 (CECR6), mRNA
NM 032863 NM 032813 NM 032578 NM 032385 NM 032239 NM 032012 NM 031922 NM 031890	Homo sapiens chromosome X open reading frame 12 (CXorf12), mRNA Homo sapiens embryonic ectoderm development (EED), mRNA Homo sapiens hypothetical protein FLJ14927 (FLJ14927), mRNA Homo sapiens hypothetical protein FLJ14624 (FLJ14624), mRNA Homo sapiens myopalladin (FLJ14437), mRNA Homo sapiens chromosome 5 open reading frame 4 (C5orf4), mRNA Homo sapiens hypothetical protein FLJ23511 (FLJ23511), mRNA Homo sapiens chromosome 9 open reading frame 5 (C9orf5), mRNA Homo sapiens RALBP1 protein (LOC83859), mRNA Homo sapiens cat eye syndrome chromosome region, candidate 6 (CECR6), mRNA Homo sapiens chromosome 17 open reading frame 1A (C17orf1A), mRNA
NM 032863 NM 032813 NM 032578 NM 032385 NM 032239 NM 032012 NM 031922 NM 031890 NM 031456	Homo sapiens chromosome X open reading frame 12 (CXorf12), mRNA Homo sapiens embryonic ectoderm development (EED), mRNA Homo sapiens hypothetical protein FLJ14927 (FLJ14927), mRNA Homo sapiens hypothetical protein FLJ14624 (FLJ14624), mRNA Homo sapiens myopalladin (FLJ14437), mRNA Homo sapiens chromosome 5 open reading frame 4 (C5orf4), mRNA Homo sapiens hypothetical protein FLJ23511 (FLJ23511), mRNA Homo sapiens chromosome 9 open reading frame 5 (C9orf5), mRNA Homo sapiens RALBP1 protein (LOC83859), mRNA Homo sapiens cat eye syndrome chromosome region, candidate 6 (CECR6), mRNA Homo sapiens chromosome 17 open reading frame 1A (C17orf1A), mRNA Homo sapiens chromosome 15 open reading frame 5 (C15orf5), mRNA
NM 032863 NM 032813 NM 032578 NM 032385 NM 032239 NM 032012 NM 031922 NM 031890 NM 031456 NM 030944	Homo sapiens chromosome X open reading frame 12 (CXorf12), mRNA Homo sapiens embryonic ectoderm development (EED), mRNA Homo sapiens hypothetical protein FLJ14927 (FLJ14927), mRNA Homo sapiens hypothetical protein FLJ14624 (FLJ14624), mRNA Homo sapiens myopalladin (FLJ14437), mRNA Homo sapiens chromosome 5 open reading frame 4 (C5orf4), mRNA Homo sapiens hypothetical protein FLJ23511 (FLJ23511), mRNA Homo sapiens chromosome 9 open reading frame 5 (C9orf5), mRNA Homo sapiens RALBP1 protein (LOC83859), mRNA Homo sapiens cat eye syndrome chromosome region, candidate 6 (CECR6), mRNA Homo sapiens chromosome 17 open reading frame 1A (C17orf1A), mRNA Homo sapiens chromosome 15 open reading frame 5 (C15orf5), mRNA Homo sapiens chromosome 1 open reading frame 21 (C1orf21), mRNA
NM 032863 NM 032813 NM 032578 NM 032385 NM 032239 NM 032012 NM 031922 NM 031890 NM 031456 NM 030944 NM 030806	Homo sapiens chromosome X open reading frame 12 (CXorf12), mRNA Homo sapiens embryonic ectoderm development (EED), mRNA Homo sapiens hypothetical protein FLJ14927 (FLJ14927), mRNA Homo sapiens hypothetical protein FLJ14624 (FLJ14624), mRNA Homo sapiens myopalladin (FLJ14437), mRNA Homo sapiens chromosome 5 open reading frame 4 (C5orf4), mRNA Homo sapiens hypothetical protein FLJ23511 (FLJ23511), mRNA Homo sapiens chromosome 9 open reading frame 5 (C9orf5), mRNA Homo sapiens RALBP1 protein (LOC83859), mRNA Homo sapiens cat eye syndrome chromosome region, candidate 6 (CECR6), mRNA Homo sapiens chromosome 17 open reading frame 1A (C17orf1A), mRNA Homo sapiens chromosome 15 open reading frame 5 (C15orf5), mRNA Homo sapiens chromosome 1 open reading frame 21 (C1orf21), mRNA Homo sapiens hypothetical protein CDA08 (CDA08), mRNA
NM 032863 NM 032813 NM 032578 NM 032385 NM 032239 NM 032012 NM 031922 NM 031890 NM 031456 NM 030944 NM 030806 NM 030790	Homo sapiens chromosome X open reading frame 12 (CXorf12), mRNA Homo sapiens embryonic ectoderm development (EED), mRNA Homo sapiens hypothetical protein FLJ14927 (FLJ14927), mRNA Homo sapiens hypothetical protein FLJ14624 (FLJ14624), mRNA Homo sapiens myopalladin (FLJ14437), mRNA Homo sapiens chromosome 5 open reading frame 4 (C5orf4), mRNA Homo sapiens hypothetical protein FLJ23511 (FLJ23511), mRNA Homo sapiens chromosome 9 open reading frame 5 (C9orf5), mRNA Homo sapiens RALBP1 protein (LOC83859), mRNA Homo sapiens cat eye syndrome chromosome region, candidate 6 (CECR6), mRNA Homo sapiens chromosome 17 open reading frame 1A (C17orf1A), mRNA Homo sapiens chromosome 15 open reading frame 5 (C15orf5), mRNA Homo sapiens chromosome 1 open reading frame 21 (C1orf21), mRNA Homo sapiens hypothetical protein CDA08 (CDA08), mRNA Homo sapiens chromosome 11 open reading frame 23 (C11orf23), mRNA
NM 032863 NM 032813 NM 032578 NM 032385 NM 032239 NM 032012 NM 031922 NM 031890 NM 031456 NM 030944 NM 030806 NM 030790 NM 018312	Homo sapiens chromosome X open reading frame 12 (CXorf12), mRNA Homo sapiens embryonic ectoderm development (EED), mRNA Homo sapiens hypothetical protein FLJ14927 (FLJ14927), mRNA Homo sapiens hypothetical protein FLJ14624 (FLJ14624), mRNA Homo sapiens myopalladin (FLJ14437), mRNA Homo sapiens chromosome 5 open reading frame 4 (C5orf4), mRNA Homo sapiens hypothetical protein FLJ23511 (FLJ23511), mRNA Homo sapiens chromosome 9 open reading frame 5 (C9orf5), mRNA Homo sapiens RALBP1 protein (LOC83859), mRNA Homo sapiens cat eye syndrome chromosome region, candidate 6 (CECR6), mRNA Homo sapiens chromosome 17 open reading frame 1A (C17orf1A), mRNA Homo sapiens chromosome 15 open reading frame 2 (C1orf21), mRNA Homo sapiens chromosome 1 open reading frame 21 (C1orf21), mRNA Homo sapiens chromosome 11 open reading frame 23 (C11orf23), mRNA Homo sapiens chromosome 11 open reading frame 23 (C11orf23), mRNA Homo sapiens malignant cell expression-enhanced gene/tumor progression-
NM 032863 NM 032813 NM 032578 NM 032385 NM 032239 NM 032012 NM 031922 NM 031890 NM 031456 NM 030944 NM 030806 NM 030790 NM 018312	Homo sapiens chromosome X open reading frame 12 (CXorf12), mRNA Homo sapiens embryonic ectoderm development (EED), mRNA Homo sapiens hypothetical protein FLJ14927 (FLJ14927), mRNA Homo sapiens hypothetical protein FLJ14624 (FLJ14624), mRNA Homo sapiens myopalladin (FLJ14437), mRNA Homo sapiens chromosome 5 open reading frame 4 (C5orf4), mRNA Homo sapiens hypothetical protein FLJ23511 (FLJ23511), mRNA Homo sapiens chromosome 9 open reading frame 5 (C9orf5), mRNA Homo sapiens RALBP1 protein (LOC83859), mRNA Homo sapiens cat eye syndrome chromosome region, candidate 6 (CECR6), mRNA Homo sapiens chromosome 17 open reading frame 1A (C17orf1A), mRNA Homo sapiens chromosome 15 open reading frame 5 (C15orf5), mRNA Homo sapiens chromosome 1 open reading frame 21 (C1orf21), mRNA Homo sapiens hypothetical protein CDA08 (CDA08), mRNA Homo sapiens chromosome 11 open reading frame 23 (C11orf23), mRNA

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NM_022163	Homo sapiens chromosome 15 open reading frame 4 (C15orf4), mRNA
NM_022107	Homo sapiens chromosome 6 open reading frame 9 (C6orf9), mRNA
NM_006781	Homo sapiens chromosome 6 open reading frame 10 (C6orf10), mRNA
NM_019895	Homo sapiens chromosome 3 open reading frame 4 (C3orf4), mRNA
NM_012265	Homo sapiens chromosome 22 open reading frame 3 (C22orf3), mRNA
NM_021254	Homo sapiens chromosome 21 open reading frame 59 (C21orf59), mRNA
NM_020645	Homo sapiens chromosome 11 open reading frame 14 (C11orf14), mRNA
NM_012112	Homo sapiens chromosome 20 open reading frame 1 (C20orf1), mRNA
NM_018555	Homo sapiens zinc finger protein 331; zinc finger protein 463 (ZNF361), mRNA
NM_019106	Homo sapiens septin 3 (SEPT3), mRNA
NM_020375	Homo sapiens chromosome 12 open reading frame 5 (C12orf5), mRNA
NM_020374	Homo sapiens chromosome 12 open reading frame 4 (C12orf4), mRNA
NM_020373	Homo sapiens chromosome 12 open reading frame 3 (C12orf3), mRNA
NM_020367	Homo sapiens chromosome 12 open reading frame 6 (C12orf6), mRNA
NM_020130	Homo sapiens chromosome 8 open reading frame 4 (C8orf4), mRNA
NM_019596	Homo sapiens chromosome 21 open reading frame 62 (C21orf62), mRNA
NM_019063	Homo sapiens chromosome 2 open reading frame 2 (C2orf2), mRNA
NM_018956	Homo sapiens chromosome 9 open reading frame 9 (C9orf9), mRNA
NM_017586	Homo sapiens chromosome 9 open reading frame 7 (C9orf7), mRNA
NM_018691	Homo sapiens chromosome 5 open reading frame 3 (C5orf3), mRNA
NM_006134	Homo sapiens chromosome 21 open reading frame 4 (C21orf4), mRNA
NM_016940	Homo sapiens chromosome 21 open reading frame 6 (C21orf6), mRNA
NM_017438	Homo sapiens chromosome 21 open reading frame 18 (C21orf18), mRNA
NM_013265	Homo sapiens chromosome 11 open reading frame2 (C11orf2), mRNA
NM_016190	Homo sapiens chromosome 1 open reading frame 10 (C1orf10), mRNA
NM_015927	Homo sapiens transforming growth factor beta 1 induced transcript 1
	(TGFB1I1), mRNA
NM_016564	Homo sapiens BM88 antigen (BM88), mRNA
NM_016348	Homo sapiens chromosome 5 open reading frame 4 (C5orf4), mRNA
NM_014009	Homo sapiens immune dysregulation, polyendocrinopathy, enteropathy, X-linked (IPEX), mRNA
NM 015524	Homo sapiens chromosome 6 open reading frame 5 (C6orf5), mRNA
NM 006345	Homo sapiens chromosome 4 open reading frame 1 (C4orf1), mRNA
NM 015373	Homo sapiens chromosome 22 open reading frame 2 (C22orf2), mRNA
NM 014205	Homo sapiens chromosome 11 open reading frame 5 (C11orf5), mRNA
NM 012264	Homo sapiens chromosome 22 open reading frame 5 (C22orf5), mRNA
NM 012111	Homo sapiens chromosome 14 open reading frame 3 (C14orf3), mRNA
NM_007211	Homo sapiens chromosome 12 open reading frame 2 (C12orf2), mRNA
NM_007176	Homo sapiens chromosome 14 open reading frame 1 (C14orf1), mRNA
NM 006706	Homo sapiens TATA box binding protein (TBP)-associated factor, RNA
_	polymerase II, S, 150kD (TAF2S), mRNA
NM 006382	Homo sapiens chromosome 17 open reading frame 1A (C17orf1A), mRNA
NM_005967	Homo sapiens NGFI-A binding protein 2 (EGR1 binding protein 2) (NAB2),
-	mRNA
NM_005966	Homo sapiens NGFI-A binding protein 1 (EGR1 binding protein 1) (NAB1), mRNA
NM 005663	Homo sapiens Wolf-Hirschhorn syndrome candidate 2 (WHSC2), mRNA
NM 005491	Homo sapiens chromosome X open reading frame 6 (CXorf6), mRNA
NM 005128	Homo sapiens chromosome 21 open reading frame 5 (C21orf5), mRNA
NM_004928	Homo sapiens chromosome 21 open reading frame 2 (C21orf2), mRNA
NM_004894	Homo sapiens chromosome 14 open reading frame 2 (C210f12), mRNA
NM_004872	Homo sapiens chromosome 1 open reading frame 8 (C1orf8), mRNA
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NM_004709	Homo sapiens chromosome X open reading frame 1 (CXorf1), mRNA
NM_004337	Homo sapiens chromosome 8 open reading frame 1 (C8orf1), mRNA
NM_004913	Homo sapiens chromosome 16 open reading frame 7 (C16orf7), mRNA
NM_000956	Homo sapiens prostaglandin E receptor 2 (subtype EP2), 53kD (PTGER2), mRNA
NM 001586	Homo sapiens chromosome X open reading frame 2 (CXorf2), mRNA
NM 001585	Homo sapiens chromosome 22 open reading frame 1 (C22orf1), mRNA
NM 001214	Homo sapiens chromosome 16 open reading frame 3 (C16orf3), mRNA
NM 001584	Homo sapiens chromosome 11 open reading frame 8 (C11orf8), mRNA
NM 003475	Homo sapiens chromosome 11 open reading frame 13 (C11orf13), mRNA
NM 032496	Homo sapiens rho-gtpase activating protein ARHGAP9 (ARHGAP9), mRNA
NM 007234	Homo sapiens dynactin 3 (p22) (DCTN3), transcript variant 1, mRNA
NM_024348	Homo sapiens dynactin 3 (p22) (DCTN3), transcript variant 2, mRNA
NM_021246	Homo sapiens megakaryocyte-enhanced gene transcript 1 protein (MEGT1), mRNA
NM_013291	Homo sapiens cleavage and polyadenylation specific factor 1, 160kD subunit (CPSF1), mRNA
NM 014500	Homo sapiens HIV TAT specific factor 1 (HTATSF1), mRNA
NM_005567	Homo sapiens lectin, galactoside-binding, soluble, 3 binding protein
NIM 005711	(LGALS3BP), mRNA
NM_005711 NM_016593	Homo sapiens EGF-like repeats and discoidin I-like domains 3 (EDIL3), mRNA
NM 021048	Homo sapiens oxysterol 7alpha-hydroxylase (CYP39A1), mRNA
NM 021049	Homo sapiens melanoma antigen, family A, 10 (MAGEA10), mRNA
NM 019602	Homo sapiens melanoma antigen, family A, 5 (MAGEA5), mRNA
	Homo sapiens butyrophilin-like 2 (MHC class II associated) (BTNL2), mRNA
NM_018002	Homo sapiens oxidation resistance 1 (OXR1), mRNA
NM_013392	Homo sapiens nuclear receptor binding protein (NRBP), mRNA
NM_012396	Homo sapiens pleckstrin homology-like domain, family A, member 3 (PHLDA3), mRNA
NM_006492	Homo sapiens aristaless-like homeobox 3 (ALX3), mRNA
NM_005365	Homo sapiens melanoma antigen, family A, 9 (MAGEA9), mRNA
NM_005364	Homo sapiens melanoma antigen, family A, 8 (MAGEA8), mRNA
NM_005366	Homo sapiens melanoma antigen, family A, 11 (MAGEA11), mRNA
NM_024490	Homo sapiens ATPase, Class V, type 10C (ATP10C), mRNA
NM_020354	Homo sapiens lysosomal apyrase-like protein 1 (LALP1), mRNA
NM_018655	Homo sapiens lens epithelial protein (LENEP), mRNA
NM_016448	Homo sapiens RA-regulated nuclear matrix-associated protein (RAMP), mRNA
NM_014763	Homo sapiens mitochondrial ribosomal protein L19 (MRPL19), mRNA
NM_006099	Homo sapiens protein inhibitor of activated STAT3 (PIAS3), mRNA
NM_004221	Homo sapiens natural killer cell transcript 4 (NK4), mRNA
NM_002949	Homo sapiens mitochondrial ribosomal protein L12 (MRPL12), mRNA
NM_016239	Homo sapiens myosin XVA (MYO15A), mRNA
NM_005094	Homo sapiens solute carrier family 27 (fatty acid transporter), member 4 (SLC27A4), mRNA
NM_015077	Homo sapiens sterile alpha and HEAT/Armadillo motif protein, ortholog of
-	Drosophila (SARM), mRNA
NM_013239	Homo sapiens protein phosphatase 2A 48 kDa regulatory subunit (PR48), mRNA
NM_022363	Homo sapiens LIM homeobox protein 5 (LHX5), mRNA
NM_031966	Homo sapiens cyclin B1 (CCNB1), mRNA
NM_015559	Homo sapiens SET binding protein 1 (SETBP1), mRNA
NM_007178	Homo sapiens unr-interacting protein (UNRIP), mRNA
NM_005367	Homo sapiens melanoma antigen, family A, 12 (MAGEA12), mRNA

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NM_031275	Homo sapiens testis expressed sequence 12 (TEX12), mRNA
NM_032403	Homo sapiens protocadherin gamma subfamily C, 3 (PCDHGC3), transcript
	variant 3, mRNA
NM_032402	Homo sapiens protocadherin gamma subfamily C, 3 (PCDHGC3), transcript
	variant 2, mRNA
NM_002588	Homo sapiens protocadherin gamma subfamily C, 3 (PCDHGC3), transcript
	variant 1, mRNA
NM_014583	Homo sapiens LIM and cysteine-rich domains 1 (LMCD1), mRNA
NM_001389	Homo sapiens Down syndrome cell adhesion molecule (DSCAM), mRNA
NM 031894	Homo sapiens ferritin, heavy polypeptide-like 17 (FTHL17), mRNA
NM 032098	Homo sapiens protocadherin gamma subfamily B, 4 (PCDHGB4), transcript
_	variant 2, mRNA
NM_003736	Homo sapiens protocadherin gamma subfamily B, 4 (PCDHGB4), transcript
-	variant 1, mRNA
NM 032938	Homo sapiens G protein pathway suppressor 2 (GPS2), transcript variant 3,
	mRNA
NM 004489	Homo sapiens G protein pathway suppressor 2 (GPS2), transcript variant 2,
	mRNA
NM 032442	Homo sapiens G protein pathway suppressor 2.(GPS2), transcript variant 1,
	mRNA
NM 001887	Homo sapiens crystallin, beta B1 (CRYBB1), mRNA
NM 005208	Homo sapiens crystallin, beta A1 (CRYBA1), mRNA
NM 001889	Homo sapiens crystallin, zeta (quinone reductase) (CRYZ), mRNA
NM_022132	Homo soniens methylenotonoul Community (CRYZ), mRNA
14141_022132	Homo sapiens methylcrotonoyl-Coenzyme A carboxylase 2 (beta) (MCCC2), mRNA
NM_001288	
NM_021624	Homo sapiens chloride intracellular channel 1 (CLIC1), mRNA
NM 032527	Homo sapiens histamine H4 receptor (HRH4), mRNA
	Homo sapiens hypothetical protein FLJ14972 (KIAA1847), mRNA
NM 005560	Homo sapiens laminin, alpha 5 (LAMA5), mRNA
NM_032931	Homo sapiens hypothetical protein MGC13219 (MGC13219), mRNA
NM 032924	Homo sapiens hypothetical protein MGC16040 (MGC16040), mRNA
NM_032920	Homo sapiens hypothetical protein MGC15873 (MGC15873), mRNA
NM 032913	Homo sapiens hypothetical protein MGC14458 (MGC14458), mRNA
NM_032893	Homo sapiens hypothetical protein MGC14336 (MGC14336), mRNA
NM_032889	Homo sapiens hypothetical protein MGC11308 (MGC11308), mRNA
NM_032815	Homo sapiens hypothetical protein FLJ14639 (FLJ14639), mRNA
NM_032798	Homo sapiens hypothetical protein FLJ14503 (FLJ14503), mRNA
NM_032793	Homo sapiens hypothetical protein FLJ14490 (FLJ14490), mRNA
NM_032791	Homo sapiens hypothetical protein FLJ14477 (FLJ14477), mRNA
NM_032789	Homo sapiens hypothetical protein FLJ14464 (FLJ14464), mRNA
NM_032769	Homo sapiens hypothetical protein MGC16212 (MGC16212), mRNA
NM_032760	Homo sapiens hypothetical protein MGC14966 (MGC14966), mRNA
NM_032696	Homo sapiens hypothetical protein MGC12262 (MGC12262), mRNA
NM_032665	Homo sapiens hypothetical protein MGC4640 (MGC4640), mRNA
NM_032662	Homo sapiens hypothetical protein MGC10600 (MGC10600), mRNA
NM_032655	Homo sapiens hypothetical protein MGC10997 (MGC10997), mRNA
NM_032625	Homo sapiens hypothetical brain protein my040 (MY040), mRNA
NM_032621	Homo sapiens X-linked protein (DJ79P11.1), mRNA
NM_032525	Homo sapiens tubulin beta-5 (TUBB5), mRNA
NM 005485	Homo sapiens ADP-ribosyltransferase (NAD+; poly (ADP-ribose) polymerase)-
_	like 3 (ADPRTL3), mRNA
NM 005484	Homo sapiens ADP-ribosyltransferase (NAD+; poly(ADP-ribose) polymerase)-
	Processia and the contract (TALDT, poly(ADT-Tiouse) polymerase)-

	like 2 (ADPRTL2), mRNA
NM_005447	Homo sapiens peptidylglycine alpha-amidating monooxygenase COOH-terminal
	interactor (PAMCI), mRNA
NM_000137	Homo sapiens fumarylacetoacetate hydrolase (fumarylacetoacetase) (FAH), mRNA
NM_001888	Homo sapiens crystallin, mu (CRYM), mRNA
NM_032608	Homo sapiens hypothetical protein bk125H2.1 (BK125H2.1), mRNA
NM_032607	Homo sapiens CREB/ATF family transcription factor (CREB-H), mRNA
NM 032602	Homo sapiens connexin 62 (CX62), mRNA
NM 032598	Homo sapiens testes development-related NYD-SP20 (NYD-SP20), mRNA
NM 032592	Homo sapiens 1-aminocyclopropane-1-carboxylate synthase (PHACS), mRNA
NM 032581	Homo sapiens down-regulated by Ctnnb1, a (DRCTNNB1A), mRNA
NM 032579	Homo sapiens colon and small intestine-specific cysteine-rich protein precursor
	similar to FIZZ2/resistin-like protein (HXCP2), mRNA
NM 032570	Homo sapiens NPC-related protein NAG73 (NAG73), mRNA
NM_032565	Homo sapiens emopamil binding related protein, delta8-delta7 sterol isomerase
	related protein (EBRP), mRNA
NM 032561	Homo sapiens EVG1 protein (EVG1), mRNA
NM 032555	Homo sapiens P143 protein (P143), mRNA
NM_032549	Homo sapiens inner mitochondrial membrane peptidase 2 like (IMMP2L),
	mRNA
NM 032548	Homo sapiens BPOZ protein (BPOZ), mRNA
NM 015080	Homo sapiens neurexin 2 (NRXN2), mRNA
NM 005676	Homo sapiens RNA binding motif protein 10 (RBM10), mRNA
NM 032526	Homo sapiens cytosolic nucleotidase I (CN-I), mRNA
NM 032483	Homo sapiens HTPAP protein (HTPAP), mRNA
NM 032094	Homo sapiens protocadherin gamma subfamily A, 12 (PCDHGA12), transcript
	variant 2, mRNA
NM_003735	Homo sapiens protocadherin gamma subfamily A, 12 (PCDHGA12), transcript variant 1, mRNA
NM_031887	Homo sapiens pro-melanin-concentrating hormone-like 1 (PMCHL1), mRNA
NM_032461	Homo sapiens SPANX family, member B1 (SPANXB1), mRNA
NM_006986	Homo sapiens melanoma antigen, family D, 1 (MAGED1), mRNA
NM_005462	Homo sapiens melanoma antigen, family C, 1 (MAGEC1), mRNA
NM_002375	Homo sapiens microtubule-associated protein 4 (MAP4), transcript variant 1,
	mRNA
NM_030983	Homo sapiens microtubule-associated protein 4 (MAP4), transcript variant 4, mRNA
NM_030885	Homo sapiens microtubule-associated protein 4 (MAP4), transcript variant 3, mRNA
NM_030884	Homo sapiens microtubule-associated protein 4 (MAP4), transcript variant 2, mRNA
NM_002374	Homo sapiens microtubule-associated protein 2 (MAP2), transcript variant 1, mRNA
NM_031847	Homo sapiens microtubule-associated protein 2 (MAP2), transcript variant 4, mRNA
NM_031846	Homo sapiens microtubule-associated protein 2 (MAP2), transcript variant 3, mRNA
NM_031845	Homo sapiens microtubule-associated protein 2 (MAP2), transcript variant 2.
NR (600 () 5	mRNA
NM_032446	Homo sapiens MEGF10 protein (MEGF10), mRNA
NM_032417	Homo sapiens SPANX family, member D (SPANXD), mRNA

NM 020690 Homo NM_012121 Homo nRNA NM 001019 Homo NM_022551 Homo	er A1 (SPANXA1), mRNA sapiens KIAA1085 protein (KIAA1085), mRNA sapiens Cdc42 effector protein 4; binder of Rho GTPases 4 (CEP4), A sapiens ribosomal protein S15a (RPS15A), mRNA sapiens ribosomal protein S18 (RPS18), mRNA sapiens microtubule-associated protein 1B (MAP1B), transcript variant 1,
NM_012121 Homo mRN/n NM_001019 Homo NM_022551	sapiens Cdc42 effector protein 4; binder of Rho GTPases 4 (CEP4), A sapiens ribosomal protein S15a (RPS15A), mRNA sapiens ribosomal protein S18 (RPS18), mRNA
mRNA NM 001019 Homo NM_022551 Homo	sapiens ribosomal protein S15a (RPS15A), mRNA sapiens ribosomal protein S18 (RPS18), mRNA
NM_022551 Homo	sapiens ribosomal protein S18 (RPS18), mRNA
	saniens microtubule-associated protein 1B (MAP1B), transcript variant 1
mRN	4
NM_032010 Homo mRNA	sapiens microtubule-associated protein 1B (MAP1B), transcript variant 2,
	sapiens microtubule-associated protein 1A (MAP1A), mRNA
transc	sapiens collagen, type IV, alpha 3 (Goodpasture antigen) (COL4A3), ript variant 6, mRNA
	sapiens collagen, type IV, alpha 3 (Goodpasture antigen) (COL4A3), ript variant 5, mRNA
	sapiens collagen, type IV, alpha 3 (Goodpasture antigen) (COL4A3), ript variant 4, mRNA
	sapiens collagen, type IV, alpha 3 (Goodpasture antigen) (COL4A3), ript variant 3, mRNA
	sapiens collagen, type IV, alpha 3 (Goodpasture antigen) (COL4A3), ript variant 2, mRNA
NM_000091 Homo	sapiens collagen, type IV, alpha 3 (Goodpasture antigen) (COL4A3), ript variant 1, mRNA
NM_002140 Homo	sapiens heterogeneous nuclear ribonucleoprotein K (HNRPK), transcript t 1, mRNA
NM_031263 Homo	sapiens heterogeneous nuclear ribonucleoprotein K (HNRPK), transcript t 3, mRNA
	sapiens heterogeneous nuclear ribonucleoprotein K (HNRPK), transcript t 2, mRNA
	sapiens prokineticin 1 precursor (PROK1), mRNA
	sapiens TEA domain family member 3 (TEAD3), mRNA
	sapiens DKFZP434K091 protein (PAL), mRNA
	sapiens apolipoprotein L, 4 (APOL4), mRNA
	sapiens hypothetical protein FLJ12565 (FLJ12565), mRNA
	sapiens mitochondrial ribosomal protein L20 (MRPL20), mRNA
	sapiens mitochondrial ribosomal protein L27 (MRPL27), mRNA
	sapiens mitochondrial ribosomal protein L42 (MRPL42), mRNA
	sapiens alpha-2-macroglobulin (A2M), mRNA
	sapiens mitochondrial ribosomal protein L33 (MRPL33), mRNA
	sapiens prostate differentiation factor (PLAB), mRNA
NM_000454 Homo	sapiens superoxide dismutase 1, soluble (amyotrophic lateral sclerosis 1)) (SOD1), mRNA
	sapiens small nuclear protein PRAC (PRAC), mRNA
	sapiens hypothetical protein FLJ22315 (FLJ22315), mRNA
	sapiens hypothetical protein MGC5254 (MGC5254), mRNA
	sapiens HEIL2 protein (HEIL2), mRNA
	sapiens hypothetical protein MGC14797 (MGC14797), mRNA
	sapiens hypothetical protein DKFZp547E052 (DKFZp547E052), mRNA
	sapiens hypothetical protein DKFZp586G1123 (DKFZp586G1123),
	sapiens hypothetical protein DKFZp434P144 (DKFZp434P144), mRNA

NM_032237	Homo sapiens hypothetical protein FLJ23356 (FLJ23356), mRNA
NM_032220	Homo sapiens hypothetical protein FLJ22283 (FLJ22283), mRNA
NM_032219	Homo sapiens hypothetical protein FLJ22269 (FLJ22269), mRNA
NM_032204	Homo sapiens hypothetical protein FLJ21588 (FLJ21588), mRNA
NM_032203	Homo sapiens hypothetical protein FLJ21423 (FLJ21423), mRNA
NM_032202	Homo sapiens hypothetical protein FLJ21404 (FLJ21404), mRNA
NM_032173	Homo sapiens hypothetical protein FLJ12747 (FLJ12747), mRNA
NM_032157	Homo sapiens hypothetical protein FLJ11531 (FLJ11531), mRNA
NM_032150	Homo sapiens hypothetical protein DKFZp434P1735 (DKFZP434P1735), mRNA
NM_021005	Homo sapiens nuclear receptor subfamily 2, group F, member 2 (NR2F2), mRNA
NM_020159	Homo sapiens hypothetical protein DKFZp762K2015 (DKFZp762K2015), mRNA
NM_015449	Homo sapiens DKFZP586G1722 protein (DKFZP586G1722), mRNA
NM_015424	Homo sapiens DKFZP586N2124 protein (DKFZP586N2124), mRNA
NM_015235	Homo sapiens likely ortholog of mouse variant polyadenylation protein CSTF-
	64; KIAA0689 protein (KIAA0689), mRNA
NM_015068	Homo sapiens paternally expressed 10 (PEG10), mRNA
NM_014599	Homo sapiens EH-domain containing 4 (EHD4), mRNA
NM_014411	Homo sapiens brain and nasopharyngeal carcinoma susceptibility protein (NSG-X), mRNA
NM_007148	Homo sapiens zinc finger protein 179 (ZNF179), mRNA
NM_007266	Homo sapiens XPA binding protein 1; putative ATP(GTP)-binding protein (NTPBP), mRNA
NM 006313	Homo sapiens ubiquitin specific protease 15 (USP15), mRNA
NM 005726	Homo sapiens Ts translation elongation factor, mitochondrial (TSFM), mRNA
NM 005277	Homo sapiens glycoprotein M6A (GPM6A), mRNA
NM 005437	Homo sapiens nuclear receptor coactivator 4 (NCOA4), mRNA
NM 001439	Homo sapiens exostoses (multiple)-like 2 (EXTL2), mRNA
NM 001287	Homo sapiens chloride channel 7 (CLCN7), mRNA
NM_021194	Homo sapiens solute carrier family 30 (zinc transporter), member 1 (SLC30A1), mRNA
NM_013986	Homo sapiens Ewing sarcoma breakpoint region 1 (EWSR1), transcript variant EWS-b, mRNA
NM 001013	Homo sapiens ribosomal protein S9 (RPS9), mRNA
NM 005617	Homo sapiens ribosomal protein S14 (RPS14), mRNA
NM 006361	Homo sapiens homeo box B13 (HOXB13), mRNA
NM 000990	Homo sapiens ribosomal protein L27a (RPL27A), mRNA
NM 005821	Homo sapiens NBR2 (NBR2), mRNA
NM_003483	Homo sapiens high-mobility group (nonhistone chromosomal) protein isoform I-C (HMGIC), mRNA
NM_002129	Homo sapiens high-mobility group (nonhistone chromosomal) protein 2 (HMG2), mRNA
NM 005959	Homo sapiens melatonin receptor 1B (MTNR1B), mRNA
NM 005958	Homo sapiens melatonin receptor 1A (MTNR1A), mRNA
NM 004739	Homo sapiens metastasis-associated 1-like 1 (MTA1L1), mRNA
NM_021644	Homo sapiens heterogeneous nuclear ribonucleoprotein H3 (2H9) (HNRPH3),
	transcript variant 2H9A, mRNA
NM_012207	Homo sapiens heterogeneous nuclear ribonucleoprotein H3 (2H9) (HNRPH3), transcript variant 2H9, mRNA
NM 019597	Homo sapiens heterogeneous nuclear ribonucleoprotein H2 (H') (HNRPH2),
1,1,1,1,01,00,1	The state of the s

	mRNA
NM_031203	
1111_031203	Homo sapiens heterogeneous nuclear ribonucleoprotein M (HNRPM), transcript variant 2, mRNA
NM_005968	
1111_005500	Homo sapiens heterogeneous nuclear ribonucleoprotein M (HNRPM), transcript variant 1, mRNA
NM 004966	Homo sapiens heterogeneous nuclear ribonucleoprotein F (HNRPF), mRNA
NM 032093	Homo sapiens pregnancy-associated interferon (HTIFN), mRNA
NM 020236	Homo seriens mitochondrial ribesonal metric T. 1.0 CDP 13
NM 016050	Homo sapiens mitochondrial ribosomal protein L1 (MRPL1), mRNA
NM_005520	Homo sapiens mitochondrial ribosomal protein L11 (MRPL11), mRNA
1411_005520	Homo sapiens heterogeneous nuclear ribonucleoprotein H1 (H) (HNRPH1), mRNA
NM_002226	Homo sapiens jagged 2 (JAG2), mRNA
NM_006805	Homo sapiens heterogeneous nuclear ribonucleoprotein A0 (HNRPA0), mRNA
NM_005463	Homo sapiens heterogeneous nuclear ribonucleoprotein D-like (HNRPDL),
-	transcript variant 1, mRNA
NM_031372	Homo sapiens heterogeneous nuclear ribonucleoprotein D-like (HNRPDL),
	transcript variant 2, mRNA
NM_031313	Homo sapiens alkaline phosphatase, placental-like 2 (ALPPL2), mRNA
NM_005080	Homo sapiens X-box binding protein 1 (XBP1), mRNA
NM_031267	Homo sapiens cell division cycle 2-like 5 (cholinesterase-related cell division
	controller) (CDC2L5), transcript variant 2, mRNA
NM_003718	Homo sapiens cell division cycle 2-like 5 (cholinesterase-related cell division
	controller) (CDC2L5), transcript variant 1, mRNA
NM_000106	Homo sapiens cytochrome P450, subfamily IID (debrisoquine, sparteine, etc., -
	metabolizing), polypeptide 6 (CYP2D6), mRNA
NM_031862	Homo sapiens membrane component, chromosome 17, surface marker 2 (ovarian
30 6 001060	carcinoma antigen CA125) (M17S2), transcript variant 3, mRNA
NM_031858	Homo sapiens membrane component, chromosome 17, surface marker 2 (ovarian
ND / 005000	carcinoma antigen CA125) (M17S2), transcript variant 2, mRNA
NM_005899	Homo sapiens membrane component, chromosome 17, surface marker 2 (ovarian
NM_032018	carcinoma antigen CA125) (M17S2), transcript variant 1, mRNA
NM_014469	Homo sapiens hypothetical protein DKFZp547N043 (DKFZP547N043), mRNA
14141_014409	Homo sapiens testes-specific heterogenous nuclear ribonucleoprotein G-T (HNRNPG-T), mRNA
NM_002137	Homo sapiens heterogeneous nuclear ribonucleoprotein A2/B1 (HNRPA2B1),
	transcript variant A2, mRNA
NM_031243	Homo sapiens heterogeneous nuclear ribonucleoprotein A2/B1 (HNRPA2B1),
	transcript variant B1, mRNA
NM_031157	Homo sapiens heterogeneous nuclear ribonucleoprotein A1 (HNRPA1),
	transcript variant 2, mRNA
NM_009585	Homo sapiens angiotensin receptor 1 (AGTR1), transcript variant 2, mRNA
NM_032049	Homo sapiens angiotensin receptor 1 (AGTR1), transcript variant 5, mRNA
NM_031850	Homo sapiens angiotensin receptor 1 (AGTR1), transcript variant 4, mRNA
NM_004835	Homo sapiens angiotensin receptor 1 (AGTR1), transcript variant 3, mRNA
NM_000685	Homo sapiens angiotensin receptor 1 (AGTR1), transcript variant 1, mRNA
NM_003965	Homo sapiens chemokine (C-C motif) receptor-like 2 (CCRL2), mRNA
NM_006641	Homo sapiens chemokine (C-C motif) receptor 9 (CCR9), transcript variant B, mRNA
NM 031200	Homo sapiens chemokine (C-C motif) receptor 9 (CCR9), transcript variant A,
	mRNA
NM_031409	Homo sapiens chemokine (C-C motif) receptor 6 (CCR6), transcript variant 2,
	mRNA

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NM_004367	Homo sapiens chemokine (C-C motif) receptor 6 (CCR6), transcript variant 1, mRNA
NM 031371	Homo sapiens RBP1-like protein (BCAA), transcript variant 2, mRNA
NM 016374	Homo sapiens RBP1-like protein (BCAA), transcript variant 1, mRNA
NM_004281	Homo sapiens BCL2-associated athanogene 3 (BAG3), mRNA
NM 032048	Homo sapiens extracellular glycoprotein EMILIN-2 precursor (EMILIN-2),
	mRNA
NM_032046	Homo sapiens mosaic serine protease (MSP), mRNA
NM_032045	Homo sapiens kringle-containing transmembrane protein; kringle-coding gene
	marking the eye and the nose (KREMEN), mRNA
NM_032044	Homo sapiens regenerating gene type IV (REG-IV), mRNA
NM_032041	Homo sapiens neurocalcin delta (NCALD), mRNA
NM_032039	Homo sapiens hypothetical protein DKFZp761D0211 (DKFZP761D0211), mRNA
NM_032038	Homo sapiens spinster-like protein (LOC83985), mRNA
NM_032020	Homo sapiens hypothetical protein MGC1314 similar to fucosidase, alpha-L-1,
	tissue (MGC1314), mRNA
NM_032016	Homo sapiens hypothetical protein MGC3251 (MGC3251), mRNA
NM_000323	Homo sapiens ret proto-oncogene (multiple endocrine neoplasia and medullary
37.6.0000==	thyroid carcinoma 1, Hirschsprung disease) (RET), transcript variant 1, mRNA
NM_020975	Homo sapiens ret proto-oncogene (multiple endocrine neoplasia and medullary
NB4 020620	thyroid carcinoma 1, Hirschsprung disease) (RET), transcript variant 2, mRNA
NM_020630	Homo sapiens ret proto-oncogene (multiple endocrine neoplasia and medullary
NM_020629	thyroid carcinoma 1, Hirschsprung disease) (RET), transcript variant 4, mRNA
141VI_020629	Homo sapiens ret proto-oncogene (multiple endocrine neoplasia and medullary
NM_016817	thyroid carcinoma 1, Hirschsprung disease) (RET), transcript variant 3, mRNA
11111_010017	Homo sapiens 2'-5'-oligoadenylate synthetase 2 (69-71 kD) (OAS2), transcript variant 1, mRNA
NM_006187	
NM_002535	Homo sapiens 2'-5'-oligoadenylate synthetase 3 (100 kD) (OAS3), mRNA Homo sapiens 2'-5'-oligoadenylate synthetase 2 (69-71 kD) (OAS2), transcript
	variant 2, mRNA
NM_002342	Homo sapiens lymphotoxin beta receptor (TNFR superfamily, member 3) (LTBR), mRNA
NM 002136	
	Homo sapiens heterogeneous nuclear ribonucleoprotein A1 (HNRPA1), transcript variant 1, mRNA
NM_001885	Homo sapiens crystallin, alpha B (CRYAB), mRNA
NM 015139	Homo sapiens UDP-glucuronic acid/UDP-N-acetylgalactosamine dual
	transporter (UGTREL7), mRNA
NM_024333	Homo sapiens fibronectin type 3 and SPRY domain-containing protein (FSD1),
	MRNA
NM_017947	Homo sapiens molybdenum cofactor sulfurase (HMCS), mRNA
NM_017934	Homo sapiens pleckstrin homology domain interacting protein (PHIP) mPNA
NM_016492	Homo sapiens homolog of yeast MOG1 (MOG1), mRNA
NM_014185	Homo sapiens homolog of yeast MOG1 (MOG1), mRNA
NM_031965	Homo sapiens haspin (GSG2), mRNA
NM_031952	Homo sapiens NYD-SP16 protein (NYD-SP16), mRNA
NM_031950	Homo sapiens Ksp37 protein (KSP37) mRNA
NM_031949	Homo sapiens NYD-TSPG protein (NYD-TSPG), mRNA
NM_031945	Homo sapiens oculospanin (OCSP), mRNA
NM_031943	Homo sapiens IFP38 (IFP38), mRNA
NM_031942	Homo sapiens c-Myc target JPO1 (JPO1), mRNA
NM_031941	Homo sapiens AIE-75 binding protein protein (MCC2), mRNA

NM_031938	Homo sapiens putative b,b-carotene-9',10'-dioxygenase (B-DIOX-II), mRNA
NM_031937	Homo sapiens EBP50-PDZ interactor of 64 kD (EPI64), mRNA
NM_031921	Homo sapiens AAA-ATPase TOB3 (TOB3), mRNA
NM_031915	Homo sapiens CLLL8 protein (CLLD8), mRNA
NM_031911	Homo sapiens complement-c1q tumor necrosis factor-related protein 7 (CTRP7), mRNA
NM_031910	Homo sapiens complement-c1q tumor necrosis factor-related protein 6 (CTRP6), mRNA
NM_031909	Homo sapiens complement-c1q tumor necrosis factor-related protein 4 (CTRP4), mRNA
NM_031904	Homo sapiens hypothetical protein FKSG44 (FKSG44), mRNA
NM_031903	Homo sapiens mitochondrial ribosomal protein L32 (MRPL32), mRNA
NM_031900	Homo sapiens alanine-glyoxylate aminotransferase 2 (AGXT2), mRNA
NM_031897	Homo sapiens calcium channel, voltage-dependent, gamma subunit 6 (CACNG6), mRNA
NM_031896	Homo sapiens calcium channel, voltage-dependent, gamma subunit 7 (CACNG7), mRNA
NM_031939	Homo sapiens B29 protein (B29), mRNA
NM_031886	Homo sapiens potassium voltage-gated channel, shaker-related subfamily, member 7 (KCNA7), mRNA
NM_020992	Homo sapiens PDZ and LIM domain 1 (elfin) (PDLIM1), mRNA
NM_031407	Homo sapiens upstream regulatory element binding protein 1 (UREB1), mRNA
NM_030582	Homo sapiens collagen, type XVIII, alpha 1 (COL18A1), mRNA
NM_020390	Homo sapiens eukaryotic translation initiation factor 5A2 (EIF5A2), mRNA
NM_018980	Homo sapiens taste receptor, type 2, member 5 (TAS2R5), mRNA
NM_018417	Homo sapiens soluble adenylyl cyclase (SAC), mRNA
NM_016945	Homo sapiens taste receptor, type 2, member 16 (TAS2R16), mRNA
NM_004775	Homo sapiens UDP-Gal:betaGlcNAc beta 1,4- galactosyltransferase, polypeptide 6 (B4GALT6), mRNA
NM_003778	Homo sapiens UDP-Gal:betaGlcNAc beta 1,4- galactosyltransferase, polypeptide 4 (B4GALT4), mRNA
NM_003779	Homo sapiens UDP-Gal:betaGlcNAc beta 1,4- galactosyltransferase, polypeptide 3 (B4GALT3), mRNA
NM 001296	Homo sapiens chemokine binding protein 2 (CCBP2), mRNA
NM_001497	Homo sapiens UDP-Gal:betaGlcNAc beta 1,4- galactosyltransferase, polypeptide
	1 (B4GALT1), mRNA
NM_014451	Homo sapiens PTH-responsive osteosarcoma B1 protein (B1), mRNA
NM_031265	Homo sapiens mucin and cadherin-like (MUCDHL), transcript variant 4, mRNA
NM_031264	Homo sapiens mucin and cadherin-like (MUCDHL), transcript variant 3, mRNA
NM_017717	Homo sapiens mucin and cadherin-like (MUCDHL), transcript variant 2, mRNA
NM_021924	Homo sapiens mucin and cadherin-like (MUCDHL), transcript variant 1, mRNA
NM_019855	Homo sapiens calcium binding protein 5 (CABP5), mRNA
NM_016367	Homo sapiens calcium binding protein 3 (CABP3), mRNA
NM_031204	Homo sapiens calcium binding protein 2 (CABP2), transcript variant 2, mRNA
NM_005201	Homo sapiens chemokine (C-C motif) receptor 8 (CCR8), mRNA
NM_000786	Homo sapiens cytochrome P450, 51 (lanosterol 14-alpha-demethylase) (CYP51), mRNA
NM_030908	Homo sapiens olfactory receptor, family 2, subfamily A, member 4 (OR2A4), mRNA
NM_001009	Homo sapiens ribosomal protein S5 (RPS5), mRNA
NM_001032	Homo sapiens ribosomal protein S29 (RPS29), mRNA
NM_001014	Homo sapiens ribosomal protein S10 (RPS10), mRNA
	The state of the property of t

Homo sapiens ribosomal protein L28 (RPL28), mRNA
220 (24 B20), HMC(A
Homo sapiens cytochrome P450, subfamily XXIV (vitamin D 24-hydroxylase)
(CYP24), mitochondrial protein encoded by nuclear gene, mRNA
Homo sapiens cytochrome P450, subfamily XIX (aromatization of androgens)
(CYP19), transcript variant 2, mRNA
Homo sapiens cytochrome P450, subfamily XIX (aromatization of androgens)
(CYP19), transcript variant 1, mRNA
Homo sapiens cytochrome P450, subfamily XIB (steroid 11-beta-hydroxylase),
polypeptide 2 (CYP11B2), mitochondrial protein encoded by nuclear gene,
mRNA
Homo sapiens cytochrome P450, subfamily XVII (steroid 17-alpha-
hydroxylase), adrenal hyperplasia (CYP17), mRNA
Homo sapiens cytochrome P450, subfamily XIB (steroid 11-beta-hydroxylase),
polypeptide 1 (CYP11B1), mitochondrial protein encoded by nuclear gene,
mRNA
Homo sapiens cytochrome P450, subfamily IIIA (niphedipine oxidase),
polypeptide 4 (CYP3A4), mRNA
Homo sapiens development and differentiation enhancing factor 1 (DDEF1), mRNA
Homo sapiens calcium binding protein 2 (CABP2), transcript variant 1, mRNA
Homo sapiens xylosylprotein beta 1,4-galactosyltransferase, polypeptide 7 (galactosyltransferase I) (B4GALT7), mRNA
Home copiens extended as PASO and family ACC 1.1 and 10.4 in the ACC 1.1 and 1
Homo sapiens cytochrome P450, subfamily 46 (cholesterol 24-hydroxylase) (CYP46), mRNA
Homo sapiens cytochrome P450, subfamily XIA (cholesterol side chain
cleavage) (CYP11A), nuclear gene encoding mitochondrial protein, mRNA
Homo sapiens chemokine (C-C motif) receptor 5 (CCR5), mRNA
Homo sapiens chemokine (C-C motif) receptor 1 (CCR1), mRNA
Homo sapiens hypothetical protein similar to RNA-binding protein lark
(MGC10871), mRNA
Homo sapiens hypothetical protein DKFZp761I141 (DKFZP761I141), mRNA
Homo sapiens SH3 domain binding glutamic acid-rich protein like 2 (SH3BGRL2), mRNA
Homo sapiens calneuron 1 (CALN1), mRNA
Homo sepiens broathetical and DVDZ SCHESSEL STATES
Homo sapiens hypothetical protein DKFZp761H2024 (DKFZP761H2024), mRNA
Homo sapiens B aggressive lymphoma gene (BAL), mRNA
Homo sapiens hypothetical protein MGC4268 (MGC4268), mRNA
Homo sapiens transmembrane protein 7 (TMEM7), mRNA
Homo sapiens retbindin (RTBDN), mRNA
Homo sapiens hypothetical protein MGC12435 (MGC12435), mRNA
Homo sapiens hypothetical protein FLJ12783 (FLJ12783), mRNA
Homo sapiens GalNAc-4-sulfotransferase 2 (GALNAC4ST-2), mRNA
Homo sapiens melanoma-derived leucine zipper, extra-nuclear factor (MLZE),
mrna
Homo sapiens cat eye syndrome chromosome region, candidate 2 (CECR2), mRNA
Homo sapiens DiGeorge syndrome critical region gene DGSI; likely ortholog of
mouse expressed sequence 2 embryonic lethal (DGSI), mRNA
Homo sapiens alcohol dehydrogenase 1C (class I), gamma polypeptide
delivered delivered to (class 1), gamma polybeptide
(ADH1C), mRNA

	mRNA
NM 018833	Homo sapiens transporter 2, ATP-binding cassette, sub-family B (MDR/TAP)
1111_010055	(TAP2), transcript variant 2, mRNA
NM 000544	Homo sapiens transporter 2, ATP-binding cassette, sub-family B (MDR/TAP)
11111_000544	(TAP2), transcript variant 1, mRNA
NM 000593	Homo sapiens transporter 1, ATP-binding cassette, sub-family B (MDR/TAP)
11111_000333	(TAP1), mRNA
NM 004678	Homo sapiens variable charge, Y chromosome, 2 (VCY2), mRNA
NM 012392	Homo sapiens PEF protein with a long N-terminal hydrophobic domain (peflin)
1111_012552	(PEF), mRNA
NM 031308	Homo sapiens epiplakin 1 (EPPK1), mRNA
NM 031299	Homo sapiens hypothetical protein MGC2577 (MGC2577), mRNA
NM_012480	Homo sapiens zinc finger protein 73 (Cos12) (ZNF73), mRNA
NM_030881	Homo sapiens DEAD/H (Asp-Glu-Ala-Asp/His) box polypeptide 17 (72kD)
2.12.12_000001	(DDX17), transcript variant 2, mRNA
NM_006386	Homo sapiens DEAD/H (Asp-Glu-Ala-Asp/His) box polypeptide 17 (72kD)
_========	(DDX17), transcript variant 1, mRNA
NM_003587	Homo sapiens DEAD/H (Asp-Glu-Ala-Asp/His) box polypeptide 16 (DDX16),
_	mRNA
NM_000478	Homo sapiens alkaline phosphatase, liver/bone/kidney (ALPL), mRNA
NM_004820	Homo sapiens cytochrome P450, subfamily VIIB (oxysterol 7 alpha-
	hydroxylase), polypeptide 1 (CYP7B1), mRNA
NM_000780	Homo sapiens cytochrome P450, subfamily VIIA (cholesterol 7 alpha-
	monooxygenase), polypeptide 1 (CYP7A1), nuclear gene encoding
	mitochondrial protein, mRNA
NM_016166	Homo sapiens DEAD/H (Asp-Glu-Ala-Asp/His) box binding protein 1
	(DDXBP1), mRNA
NM_016373	Homo sapiens WW domain-containing oxidoreductase (WWOX), mRNA
NM_024164	Homo sapiens tryptase beta 2 (TPSB2), mRNA
NM_003294	Homo sapiens tryptase beta 1 (TPSB1), mRNA
NM_031310	Homo sapiens fenestrated-endothelial linked structure protein; PV-1 protein
	(PV1), mRNA
NM_031302	Homo sapiens gycosyltransferase (LOC83468), mRNA
NM_031300	Homo sapiens hypothetical protein MGC2383 (MGC2383), mRNA
NM_031297	Homo sapiens hypothetical protein DKFZp761H1710 (DKFZP761H1710),
NR 6 001007	mRNA
NM_031287	Homo sapiens hypothetical protein MGC3133 (MGC3133), mRNA
NM_031286	Homo sapiens SH3BGRL3-like protein (SH3BGRL3), mRNA
NM_031285	Homo sapiens hypothetical protein PP1057 (PP1057), mRNA
NM_031279	Homo sapiens alanine-glyoxylate aminotransferase 2-like 1 (AGXT2L1), mRNA
NM_030970	Homo sapiens hypothetical protein MGC3771 (MGC3771), mRNA
NM 014357	Homo sapiens skin-specific protein (XP5), mRNA
NM 030590	Homo sapiens matrilin 4 (MATN4), transcript variant 2, mRNA
NM_031246	Homo sapiens pregnancy specific beta-1-glycoprotein 2 (PSG2), mRNA
NM 017422	Homo sapiens calmodulin-like skin protein (CLSP), mRNA
NM_005956	Homo sapiens methylenetetrahydrofolate dehydrogenase (NADP+ dependent),
	methenyltetrahydrofolate cyclohydrolase, formyltetrahydrofolate synthetase
NIM OCCOCC	(MTHFD1), mRNA
NM 005906	Homo sapiens male germ cell-associated kinase (MAK), mRNA
NM_006389	Homo sapiens oxygen regulated protein (150kD) (ORP150), mRNA
NM 030084	Homo sapiens organic cationic transporter-like 4 (ORCTL4), mRNA
NM_030984	Homo sapiens thromboxane A synthase 1 (platelet, cytochrome P450, subfamily

	V) (TBXAS1), transcript variant TXS-II, mRNA
NM_001061	Homo sapiens thromboxane A synthase 1 (platelet, cytochrome P450, subfamily
•	V) (TBXAS1), transcript variant TXS-I, mRNA
NM_000773	Homo sapiens cytochrome P450, subfamily IIE (ethanol-inducible) (CYP2E),
	mRNA
NM_030592	Homo sapiens matrilin 4 (MATN4), transcript variant 3, mRNA
NM_003833	Homo sapiens matrilin 4 (MATN4), transcript variant 1, mRNA
NM_005355	Homo sapiens kinesin-like 3 (KNSL3), transcript variant 2, mRNA
NM_030615	Homo sapiens kinesin-like 3 (KNSL3), transcript variant 1, mRNA
NM_004523	Homo sapiens kinesin-like 1 (KNSL1), mRNA
NM_005000	Homo sapiens NADH dehydrogenase (ubiquinone) 1 alpha subcomplex, 5
	(13kD, B13) (NDUFA5), nuclear gene encoding mitochondrial protein, mRNA
NM_004541	Homo sapiens NADH dehydrogenase (ubiquinone) 1 alpha subcomplex, 1
	(7.5kD, MWFE) (NDUFA1), nuclear gene encoding mitochondrial protein,
	mRNA
NM_000771	Homo sapiens cytochrome P450, subfamily IIC (mephenytoin 4-hydroxylase),
l	polypeptide 9 (CYP2C9), mRNA
NM_000772	Homo sapiens cytochrome P450, subfamily IIC (mephenytoin 4-hydroxylase),
37.5.04	polypeptide 18 (CYP2C18), mRNA
NM_017778	Homo sapiens Wolf-Hirschhorn syndrome candidate 1-like 1 (WHSC1L1),
ND 6 000004	transcript variant short, mRNA
NM_023034	Homo sapiens Wolf-Hirschhorn syndrome candidate 1-like 1 (WHSC1L1),
37.6.0007.66	transcript variant long, mRNA
NM_000766	Homo sapiens cytochrome P450, subfamily IIA (phenobarbital-inducible),
3D4 006646	polypeptide 13 (CYP2A13), mRNA
NM_006646	Homo sapiens WAS protein family, member 3 (WASF3), mRNA
NM_018560	Homo sapiens WW domain-containing oxidoreductase (WWOX), mRNA
NM_014110	Homo sapiens protein phosphatase 1, regulatory (inhibitor) subunit 8 (PPP1R8), mRNA
NM_004109	Homo sapiens ferredoxin 1 (FDX1), nuclear gene encoding mitochondrial
	protein, mRNA
NM_030671	Homo sapiens protein tyrosine phosphatase, receptor type, O (PTPRO),
	transcript variant 5, mRNA
NM_030670	Homo sapiens protein tyrosine phosphatase, receptor type, O (PTPRO),
	transcript variant 6, mRNA
NM_030669	Homo sapiens protein tyrosine phosphatase, receptor type, O (PTPRO),
	transcript variant 3, mRNA
NM_030668	Homo sapiens protein tyrosine phosphatase, receptor type, O (PTPRO),
	transcript variant 4, mRNA
NM_030667	Homo sapiens protein tyrosine phosphatase, receptor type, O (PTPRO),
\	transcript variant 1, mRNA
NM_002848	Homo sapiens protein tyrosine phosphatase, receptor type, O (PTPRO),
3D (001070	transcript variant 2, mRNA
NM_021979	Homo sapiens heat shock 70kD protein 2 (HSPA2), mRNA
NM_024005	Homo sapiens DEAD/H (Asp-Glu-Ala-Asp/His) box polypeptide 3 (DDX3),
ND (001256	transcript variant 1, mRNA
NM_001356	Homo sapiens DEAD/H (Asp-Glu-Ala-Asp/His) box polypeptide 3 (DDX3),
ND 6 020216	transcript variant 2, mRNA
NM_020216	Homo sapiens arginyl aminopeptidase (aminopeptidase B) (RNPEP), mRNA
NM_006990	Homo sapiens WAS protein family, member 2 (WASF2), mRNA
NM_012467	Homo sapiens tryptase gamma 1 (TPSG1), mRNA
NM_007317	Homo sapiens kinesin-like 4 (KNSL4), mRNA

NM_004256	Homo sapiens organic cationic transporter-like 3 (ORCTL3), mRNA
NM_000774	Homo sapiens cytochrome P450, subfamily IIF, polypeptide 1 (CYP2F1),
_	mRNA
NM_000769	Homo sapiens cytochrome P450, subfamily IIC (mephenytoin 4-hydroxylase),
_	polypeptide 19 (CYP2C19), mRNA
NM 031220	Homo sapiens PYK2 N-terminal domain-interacting receptor 1 (NIR1), mRNA
NM 031212	Homo sapiens hypothetical protein NPD016 (NPD016), mRNA
NM 031211	Homo sapiens LAT1-3TM protein (LAT1-3TM), mRNA
NM 031209	Homo sapiens tRNA-guanine transglycosylase (TGT), mRNA
NM 031206	Homo sapiens hypothetical protein FLJ12525 (FLJ12525), mRNA
NM_006904	Homo sapiens protein kinase, DNA-activated, catalytic polypeptide (PRKDC),
_	mRNA
NM 030963	Homo sapiens hypothetical protein DKFZp434O1427 (DKFZP434O1427),
_	mRNA
NM 030931	Homo sapiens epididymal secretory protein ESP13.2 (ESP13.2), mRNA
NM 030905	Homo sapiens olfactory receptor, family 2, subfamily J, member 2 (OR2J2),
_	mRNA
NM 030903	Homo sapiens olfactory receptor, family 2, subfamily W, member 1 (OR2W1),
_	mRNA
NM_012377	Homo sapiens olfactory receptor, family 7, subfamily C, member 2 (OR7C2),
_	mRNA
NM 030981	Homo sapiens small GTP-binding protein (RAB1B), mRNA
NM_030974	Homo sapiens hypothetical protein DKFZp434N1923 (DKFZP434N1923),
_	mRNA
NM_030973	Homo sapiens hypothetical protein TCBAP0758 (TCBAP0758), mRNA
NM 030968	Homo sapiens G protein coupled receptor interacting protein, complement-c1q
	tumor necrosis factor-related (ZSIG37), mRNA
NM_030945	Homo sapiens complement-clq tumor necrosis factor-related protein; likely
	ortholog of mouse CORS26 (collagenous repeat-containing sequence of 26-kDa
	protein) (CTRP3), mRNA
NM_030936	Homo sapiens hypothetical protein DKFZp434C135 (DKFZP434C135), mRNA
NM_030935	Homo sapiens TSC-22-like (THG-1), mRNA
NM_030926	Homo sapiens integral membrane protein 3 (ITM3), mRNA
NM_030893	Homo sapiens CD1E antigen, e polypeptide (CD1E), mRNA
NM_014067	Homo sapiens LRP16 protein (LRP16), mRNA
NM_030661	Homo sapiens homeo box A3 (HOXA3), mRNA
NM_030879	Homo sapiens Small evolutionarily conserved RNA, resembling C/D box small
	nucleolar (X102), mRNA
NM_012373	Homo sapiens olfactory receptor, family 3, subfamily A, member 3 (OR3A3),
	mRNA
NM_015072	Homo sapiens KIAA0998 protein (KIAA0998), mRNA
NM_030882	Homo sapiens apolipoprotein L, 2 (APOL2), mRNA
NM_002623	Homo sapiens prefoldin 4 (PFDN4), mRNA
NM 022167	Homo sapiens xylosyltransferase II (XT2), mRNA
NM_017506	Homo sapiens olfactory receptor, family 7, subfamily C, member 1 (OR7C1),
	mRNA
NM_003372	Homo sapiens von Hippel-Lindau binding protein 1 (VBP1), mRNA
NM_016097	Homo sapiens HSPC039 protein (HSPC039), mRNA
NM_014646	Homo sapiens lipin 2 (LPIN2), mRNA
NM_005880	Homo sapiens DnaJ (Hsp40) homolog, subfamily A, member 2 (DNAJA2),
	mRNA
NM 006755	Homo sapiens transaldolase 1 (TALDO1), mRNA

NM_005137	Homo sapiens DiGeorge syndrome critical region gene 2 (DGCR2), mRNA
NM_000022	Homo sapiens adenosine deaminase (ADA), mRNA
NM_003215	Homo sapiens tec protein tyrosine kinase (TEC), mRNA
NM_018425	Homo sapiens phosphatidylinositol 4-kinase type II (PI4KII), mRNA
NM_025238	Homo sapiens BTB (POZ) domain containing 1 (BTBD1), mRNA
NM_004248_	Homo sapiens G protein-coupled receptor 10 (GPR10), mRNA
NM_001642	Homo sapiens amyloid beta (A4) precursor-like protein 2 (APLP2), mRNA
NM_030821	Homo sapiens group XII secreted phospholipase A2 (PLA2G12), mRNA
NM_030820	Homo sapiens hypothetical protein DKFZp564B052 (DKFZp564B052), mRNA
NM_030816	Homo sapiens hypothetical protein DKFZp566D1346 (DKFZP566D1346), mRNA
NM_030807	Homo sapiens glucose transporter protein 10 (GLUT10), mRNA
NM_030798	Homo sapiens hypothetical protein DKFZp434D0421 (DKFZP434D0421), mRNA
NM_030797	Homo sapiens hypothetical protein DKFZp566A1524 (DKFZP566A1524), mRNA
NM_030788	Homo sapiens DC-specific transmembrane protein (LOC81501), mRNA
NM_030787	Homo sapiens factor H-related protein 5 (FHR5), mRNA
NM_030786	Homo sapiens intermediate filament protein syncoilin (SYNCOILIN), mRNA
NM_030785	Homo sapiens ortholog of mouse radial spokehead-like 1 (RSHL1), mRNA
NM_030784	Homo sapiens brain expressed G-protein-coupled receptor PSP24 beta (PSP24B), mRNA
NM_030783	Homo sapiens phosphatidylserine synthase 2 (PTDSS2), mRNA
NM_030779	Homo sapiens Eag-related gene member 2 (ERG2), mRNA
NM_030774	Homo sapiens prostate specific G-protein coupled receptor (PSGR), mRNA
NM_030772	Homo sapiens connexin 59 (GJA10), mRNA
NM_030764	Homo sapiens SH2 domain-containing phosphatase anchor protein 1 (SPAP1), mRNA
NM 030763	Homo sapiens nucleosomal binding protein 1 (NSBP1), mRNA
NM 030757	Homo sapiens makorin, ring finger protein, 4 (MKRN4), mRNA
NM_021813	Homo sapiens BTB and CNC homology 1, basic leucine zipper transcription factor 2 (BACH2), mRNA
NM 020819	Homo sapiens KIAA1411 protein (KIAA1411), mRNA
NM_030751	Homo sapiens transcription factor 8 (represses interleukin 2 expression) (TCF8), mRNA
NM_030754	Homo sapiens serum amyloid A2 (SAA2), mRNA
NM_030752	Homo sapiens t-complex 1 (TCP1), mRNA
NM_030756	Homo sapiens transcription factor 7-like 2 (T-cell specific, HMG-box) (TCF7L2), mRNA
NM_006010	Homo sapiens arginine-rich, mutated in early stage tumors (ARMET), mRNA
NM_001182	Homo sapiens aldehyde dehydrogenase 7 family, member A1 (ALDH7A1), mRNA
NM_000382	Homo sapiens aldehyde dehydrogenase 3 family, member A2 (ALDH3A2), mRNA
NM_003486	Homo sapiens solute carrier family 7 (cationic amino acid transporter, y+system), member 5 (SLC7A5), mRNA
NM_000694	Homo sapiens aldehyde dehydrogenase 3 family, member B1 (ALDH3B1), mRNA
NM_000693	Homo sapiens aldehyde dehydrogenase 1 family, member A3 (ALDH1A3), mRNA
NM_030381	Homo sapiens GLI-Kruppel family member GLI2 (GLI2), transcript variant 3, mRNA

NM_030380	Homo sapiens GLI-Kruppel family member GLI2 (GLI2), transcript variant 2, mRNA
NM_030379	Homo sapiens GLI-Kruppel family member GLI2 (GLI2), transcript variant 1, mRNA
NM_020166	Homo sapiens methylcrotonoyl-Coenzyme A carboxylase 1 (alpha) (MCCC1), mRNA
NM_005270	Homo sapiens GLI-Kruppel family member GLI2 (GLI2), transcript variant 4, mRNA
NM 002381	Homo sapiens matrilin 3 (MATN3) precursor, mRNA
NM 030583	Homo sapiens matrilin 2 (MATN2) precursor, transcript variant 2, mRNA
NM 002380	Homo sapiens matrilin 2 (MATN2) precursor, transcript variant 1, mRNA
NM 002379	Homo sapiens matrilin 1, cartilage matrix protein (MATN1), mRNA
NM_000168	Homo sapiens GLI-Kruppel family member GLI3 (Greig cephalopolysyndactyly syndrome) (GLI3), mRNA
NM 003462	Homo sapiens dynein, axonemal, light intermediate polypeptide (P28), mRNA
NM 017493	Homo sapiens Hin-1 (HSHIN1), mRNA
NM_005602	Homo sapiens claudin 11 (oligodendrocyte transmembrane protein) (CLDN11), mRNA
NM 001195	Homo sapiens beaded filament structural protein 1, filensin (BFSP1), mRNA
NM 004987	Homo sapiens LIM and senescent cell antigen-like domains 1 (LIMS1), mRNA
NM 000412	Homo sapiens histidine-rich glycoprotein (HRG), mRNA
NM 024494	Homo sapiens wingless-type MMTV integration site family, member 2B
	(WNT2B), transcript variant WNT-2B2, mRNA
NM 004993	Homo sapiens Machado-Joseph disease (spinocerebellar ataxia 3,
	olivopontocerebellar ataxia 3, autosomal dominant, ataxin 3) (MJD), transcript
	variant 1, mRNA
NM_004185	Homo sapiens wingless-type MMTV integration site family, member 2B (WNT2B), transcript variant WNT-2B1, mRNA
NM 024415	Homo sapiens VASA protein (VASA), transcript variant 2, mRNA
NM_004398	Homo sapiens DEAD/H (Asp-Glu-Ala-Asp/His) box polypeptide 10 (RNA helicase) (DDX10), mRNA
NM_004397	Homo sapiens DEAD/H (Asp-Glu-Ala-Asp/His) box polypeptide 6 (RNA helicase, 54kD) (DDX6), mRNA
NM_004396	Homo sapiens DEAD/H (Asp-Glu-Ala-Asp/His) box polypeptide 5 (RNA helicase, 68kD) (DDX5), mRNA
NM_030588	Homo sapiens DEAD/H (Asp-Glu-Ala-Asp/His) box polypeptide 9 (RNA helicase A, nuclear DNA helicase II; leukophysin) (DDX9), transcript variant 2, mRNA
NM_001357	Homo sapiens DEAD/H (Asp-Glu-Ala-Asp/His) box polypeptide 9 (RNA helicase A, nuclear DNA helicase II; leukophysin) (DDX9), transcript variant 1, mRNA
NM_004660	Homo sapiens DEAD/H (Asp-Glu-Ala-Asp/His) box polypeptide, Y chromosome (DBY), mRNA
NM 019039	Homo sapiens VASA protein (VASA), transcript variant 1, mRNA
NM_012382	Homo sapiens osmosis responsive factor (OSRF), mRNA
NM_000387	Homo sapiens solute carrier family 25 (carnitine/acylcarnitine translocase),
-	member 20 (SLC25A20), mitochondrial protein encoded by nuclear gene,
·	mRNA
NM_007240	Homo sapiens dual specificity phosphatase 12 (DUSP12), mRNA
NM_004940	Homo sapiens DEAD/H (Asp-Glu-Ala-Asp/His) box polypeptide 7 (RNA helicase, 52kD) (DDX7), mRNA
NM_004939	Homo sapiens DEAD/H (Asp-Glu-Ala-Asp/His) box polypeptide 1 (DDX1),

<u> </u>	mRNA
NM 013366	Homo sapiens anaphase-promoting complex subunit 2 (APC2), mRNA
NM 003791	Homo sapiens membrane-bound transcription factor protease, site 1 (MBTPS1),
14tM_003731	mRNA
NM_002251	Homo sapiens potassium voltage-gated channel, delayed-rectifier, subfamily S,
1414_002251	member 1 (KCNS1), mRNA
NM 006903	Homo sapiens inorganic pyrophosphatase (SID6-306), mRNA
NM 020956	Homo sapiens periaxin (KIAA1620), mRNA
NM 015435	Homo sapiens double ring-finger protein, Dorfin (DORFIN), mRNA
NM 014338	Homo sapiens phosphatidylserine decarboxylase (PISD), mRNA
NM 021954	Homo sapiens gap junction protein, alpha 3, 46kD (connexin 46) (GJA3), mRNA
NM 023068	Homo sapiens sialoadhesin (SN), mRNA
NM_022821	Homo sapiens elongation of very long chain fatty acids (FEN1/Elo2, SUR4/Elo3,
14141_022021	yeast)-like 1 (ELOVL1), mRNA
NM 021126	Homo sapiens mercaptopyruvate sulfurtransferase (MPST), mRNA
NM 030666	Homo sapiens serine (or cysteine) proteinase inhibitor, clade B (ovalbumin),
1111_00000	member 1 (SERPINB1), mRNA
NM 024014	Homo sapiens homeo box A6 (HOXA6), mRNA
NM 030665	Homo sapiens retinoic acid induced 1 (RAI1), mRNA
NM 030663	Homo sapiens mitochondrial capsule selenoprotein (MCSP), mRNA
NM 030664	Homo sapiens phosphotriesterase related (PTER), mRNA
NM 030662	Homo sapiens mitogen-activated protein kinase kinase 2 (MAP2K2), mRNA
NM 024896	Homo sapiens hypothetical protein FLJ23309 (FLJ23309), mRNA
NM 002183	Homo sapiens interleukin 3 receptor, alpha (low affinity) (IL3RA), mRNA
NM 021244	Homo sapiens Rag D protein; hypothetical GTP-binding protein
_	DKFZp761H171 (RAGD), mRNA
NM 005088	Homo sapiens DNA segment on chromosome X and Y (unique) 155 expressed
_	sequence (DXYS155E), mRNA
NM_016090	Homo sapiens RNA binding motif protein 7 (RBM7), mRNA
NM_013306	Homo sapiens sorting nexin 15 (SNX15), mRNA
NM_018362	Homo sapiens likely ortholog of mouse LIN-7C; mammalian LIN-7 protein 3
	(LIN-7-C), mRNA
NM_018300	Homo sapiens zinc finger protein 83 (HPF1) (ZNF83), mRNA
NM_014754	Homo sapiens phosphatidylserine synthase 1 (PTDSS1), mRNA
NM_006140	Homo sapiens colony stimulating factor 2 receptor, alpha, low-affinity
	(granulocyte-macrophage) (CSF2RA), mRNA
NM_004043	Homo sapiens acetylserotonin O-methyltransferase (ASMT), mRNA
NM_002414	Homo sapiens antigen identified by monoclonal antibodies 12E7, F21 and O13
	(MIC2), mRNA
NM_002186	Homo sapiens interleukin 9 receptor (IL9R), mRNA
NM_030657	Homo sapiens lens intrinsic membrane protein 2 (19kD) (LIM2), mRNA
NM_014349	Homo sapiens apolipoprotein L, 3 (APOL3), mRNA
NM_022566	Homo sapiens mesoderm development candidate 1 (MESDC1), mRNA
NM_020727	Homo sapiens zinc finger protein 295 (ZNF295), mRNA
NM_012074	Homo sapiens cer-d4 (mouse) homolog (CERD4), mRNA
NM_000861	Homo sapiens histamine receptor H1 (HRH1), mRNA
NM_006273	Homo sapiens small inducible cytokine A7 (monocyte chemotactic protein 3)
NR (002205	(SCYA7), mRNA
NM_002395	Homo sapiens malic enzyme 1, NADP(+)-dependent, cytosolic (ME1), mRNA
NM_024165	Homo sapiens PHD finger protein 1 (PHF1), transcript variant 2, mRNA
NM_002636	Homo sapiens PHD finger protein 1 (PHF1), transcript variant 1, mRNA
NM_001082	Homo sapiens cytochrome P450, subfamily IVF, polypeptide 2 (CYP4F2),

	mRNA
NM_007253	Homo sapiens cytochrome P450, subfamily IVF, polypeptide 8 (CYP4F8),
1414_007255	mRNA
NM 000779	Homo sapiens cytochrome P450, subfamily IVB, polypeptide 1 (CYP4B1),
1111_000775	mRNA
NM 001514	Homo sapiens general transcription factor IIB (GTF2B), mRNA
NM 004127	Homo sapiens G protein pathway suppressor 1 (GPS1), mRNA
NM 024423	Homo sapiens desmocollin 3 (DSC3), transcript variant Dsc3b, mRNA
NM 001941	Homo sapiens desmocollin 3 (DSC3),transcript variant Dsc3a, mRNA
NM 004949	Homo sapiens desmocollin 2 (DSC2), transcript variant Dsc2b, mRNA
NM 024422	Homo sapiens desmocollin 2 (DSC2), transcript variant Dsc2a, mRNA
NM 004948	Homo sapiens desmocollin 1 (DSC1), transcript variant Dsc1b, mRNA
NM 024421	Homo sapiens desmocollin 1 (DSC1), transcript variant Dsc1a, mRNA
NM 001923	Homo sapiens damage-specific DNA binding protein 1 (127kD) (DDB1), mRNA
NM 000425	Homo sapiens L1 cell adhesion molecule (hydrocephalus, stenosis of aqueduct of
	Sylvius 1, MASA (mental retardation, aphasia, shuffling gait and adducted
	thumbs) syndrome, spastic paraplegia 1) (L1CAM), transcript variant 1, mRNA
NM 024003	Homo sapiens L1 cell adhesion molecule (hydrocephalus, stenosis of aqueduct of
_	Sylvius 1, MASA (mental retardation, aphasia, shuffling gait and adducted
_	thumbs) syndrome, spastic paraplegia 1) (L1CAM), transcript variant 2, mRNA
NM_004110	Homo sapiens ferredoxin reductase (FDXR), transcript variant 2, nuclear gene
	encoding mitochondrial protein, mRNA
NM_024417	Homo sapiens ferredoxin reductase (FDXR), transcript variant 1, nuclear gene
	encoding mitochondrial protein, mRNA
NM_023944	Homo sapiens cytochrome P450 isoform 4F12 (CYP4F12), mRNA
NM_022845	Homo sapiens core-binding factor, beta subunit (CBFB), transcript variant 1,
	mRNA
NM_022041	Homo sapiens giant axonal neuropathy (gigaxonin) (GAN), mRNA
NM_021187	Homo sapiens cytochrome P450, subfamily IVF, polypeptide 11 (CYP4F11),
	mRNA
NM_019599	Homo sapiens taste receptor, type 2, member 1 (TAS2R1), mRNA
NM_017579	Homo sapiens deleted in malignant brain tumors 1 (DMBT1), transcript variant
NR 015670	3, mRNA
NM 015670	Homo sapiens sentrin/SUMO-specific protease 3 (SENP3), mRNA
NM_012096	Homo sapiens adaptor protein containing pH domain, PTB domain and leucine zipper motif (APPL), mRNA
NM 005392	Homo sapiens PHD finger protein 2 (PHF2), mRNA
NM_000896	Homo sapiens cytochrome P450, subfamily IVF, polypeptide 3 (leukotriene B4
14147_000020	omega hydroxylase) (CYP4F3), mRNA
NM_022661	Homo sapiens SPANX family, member C (SPANXC), mRNA
NM_022573	Homo sapiens TSPYq1 (TSPYQ1), mRNA
NM_022089	Homo sapiens putative ATPase (HSA9947), mRNA
NM_025228	Homo sapiens hypothetical protein dJ434O14.3 (DJ434O14.3), mRNA
NM 025013	Homo sapiens KIAA1031 protein (KIAA1031), mRNA
NM 025027	Homo sapiens hypothetical protein FLJ14260 (FLJ14260), mRNA
NM 022102	Homo sapiens hypothetical protein FLJ20958 (FLJ20958), mRNA
NM 021724	Homo sapiens nuclear receptor subfamily 1, group D, member 1 (NR1D1),
	mRNA
NM 030570	Homo sapiens hypothetical protein MGC10902 (MGC10902), mRNA
NM 025135	Homo sapiens hypothetical protein FLJ22297 (KIAA1695), mRNA
NM 024317	Homo sapiens immunoglobulin-like transcript 10 (ILT10), mRNA
NM 021822	Homo sapiens phorbolin-like protein MDS019 (MDS019), mRNA

NM_017509	Homo sapiens ACO for serine protease homologue (HSRNASPH), mRNA
NM 005583	Homo sapiens lymphoblastic leukemia derived sequence 1 (LYL1), mRNA
NM 020070	Homo sapiens immunoglobulin lambda-like polypeptide 1 (IGLL1), mRNA
NM 002383	Homo sapiens MYC-associated zinc finger protein (purine-binding transcription
	factor) (MAZ), mRNA
NM 016944	Homo sapiens taste receptor, type 2, member 4 (TAS2R4), mRNA
NM 016943	Homo sapiens taste receptor, type 2, member 3 (TAS2R3), mRNA
NM 000378	Homo sapiens Wilms tumor 1 (WT1), transcript variant A, mRNA
NM 024426	Homo sapiens Wilms tumor 1 (WT1), transcript variant D, mRNA
NM 024425	Homo sapiens Wilms tumor 1 (WT1), transcript variant C, mRNA
NM 024424	Homo sapiens Wilms tumor 1 (WT1), transcript variant B, mRNA
NM 000765	Homo sapiens cytochrome P450, subfamily IIIA, polypeptide 7 (CYP3A7),
_	mRNA
NM 021570	Homo sapiens BarH-like homeobox 1 (BARX1), mRNA
NM 000068	Homo sapiens calcium channel, voltage-dependent, P/Q type, alpha 1A subunit
_	(CACNA1A), transcript variant 1, mRNA
NM_030574	Homo sapiens hypothetical protein MGC10327 (MGC10327), mRNA
NM_030573	Homo sapiens hypothetical protein MGC10963 (MGC10963), mRNA
NM_024867	Homo sapiens hypothetical protein FLJ23577 (FLJ23577), mRNA
NM_002739	Homo sapiens protein kinase C, gamma (PRKCG), mRNA
NM_020548	Homo sapiens diazepam binding inhibitor (GABA receptor modulator, acyl-
_	Coenzyme A binding protein) (DBI), mRNA
NM_025176	Homo sapiens KIAA0980 protein (KIAA0980), mRNA
NM_003789	Homo sapiens TNFRSF1A-associated via death domain (TRADD), mRNA
NM_017541	Homo sapiens crystallin, gamma S (CRYGS), mRNA
NM_006891	Homo sapiens crystallin, gamma D (CRYGD), mRNA
NM_020989	Homo sapiens crystallin, gamma C (CRYGC), mRNA
NM_005210	Homo sapiens crystallin, gamma B (CRYGB), mRNA
NM_014617	Homo sapiens crystallin, gamma A (CRYGA), mRNA
NM_002396	Homo sapiens malic enzyme 2, NAD(+)-dependent, mitochondrial (ME2),
	nuclear gene encoding mitochondrial protein, mRNA
NM_025268	Homo sapiens hypothetical protein MGC4659 (MGC4659), mRNA
NM_025244	Homo sapiens testis specific, 10 (TSGA10), mRNA
NM_025240	Homo sapiens B7 homolog 3 (B7-H3), mRNA
NM_025237	Homo sapiens sclerostin (SOST), mRNA
NM_025236	Homo sapiens HZFw1 protein (HZFW1), mRNA
NM_025235	Homo sapiens tankyrase 2 (TNKL), mRNA
NM_025233	Homo sapiens nucleotide binding protein (NBP), mRNA
NM_025232	Homo sapiens hypothetical protein FLJ22246 (FLJ22246), mRNA
NM_025218	Homo sapiens UL16-binding protein 1 (ULBP1), mRNA
NM_025217	Homo sapiens UL16-binding protein 2 (ULBP2), mRNA
NM_025215	Homo sapiens pseudouridine synthase 1 (PUS1), mRNA
NM_025214	Homo sapiens CTCL tumor antigen se57-1 (SE57-1), mRNA
NM_025212	Homo sapiens Dvl-binding protein IDAX (inhibition of the Dvl and Axin
	complex) (IDAX), mRNA
NM_025210	Homo sapiens type 1 protein phosphatase inhibitor (I-4), mRNA
NM_025209	Homo sapiens enhancer of polycomb 1 (EPC1), mRNA
NM_025205	Homo sapiens hypothetical protein DKFZp434N185 (DKFZP434N185), mRNA
NM_025198	Homo sapiens transcription termination factor-like protein (LOC80298), mRNA
NM_025193	Homo sapiens 3 beta-hydroxy-delta 5-C27-steroid oxidoreductas (C(27)-3BETA-HSD), mRNA
NM_025180	Homo sapiens hypothetical protein FLJ13386 (FLJ13386), mRNA

NM_025161	Homo sapiens hypothetical protein FLJ22175 (FLJ22175), mRNA
NM_025158	Homo sapiens hypothetical protein FLJ22251 (FLJ22251), mRNA
NM_025148	Homo sapiens hypothetical protein FLJ12986 (FLJ12986), mRNA
NM_025137	Homo sapiens hypothetical protein FLJ21439 (FLJ21439), mRNA
NM_025116	Homo sapiens hypothetical protein FLJ12781 (FLJ12781), mRNA
NM_025114	Homo sapiens hypothetical protein FLJ13615 (FLJ13615), mRNA
NM_025083	Homo sapiens hypothetical protein FLJ21128 (FLJ21128), mRNA
NM_025054	Homo sapiens hypothetical protein FLJ23132 (FLJ23132), mRNA
NM_025017	Homo sapiens hypothetical protein FLJ13892 (FLJ13892), mRNA
NM_025011	Homo sapiens hypothetical protein FLJ13744 (FLJ13744), mRNA
NM_024995	Homo sapiens hypothetical protein FLJ12616 (FLJ12616), mRNA
NM_024987	Homo sapiens hypothetical protein FLJ12345 (FLJ12345), mRNA
NM_024900	Homo sapiens hypothetical protein FLJ22479 (FLJ22479), mRNA
NM_024874	Homo sapiens hypothetical protein FLJ14225 (FLJ14225), mRNA
NM 024873	Homo sapiens hypothetical protein FLJ21162 (FLJ21162), mRNA
NM_024861	Homo sapiens hypothetical protein FLJ22671 (FLJ22671), mRNA
NM_024836	Homo sapiens hypothetical protein FLJ22301 (FLJ22301), mRNA
NM 024822	Homo sapiens hypothetical protein FLJ22601 (FLJ22601), mRNA
NM_024819	Homo sapiens hypothetical protein FLJ22955 (FLJ22955), mRNA
NM_024816	Homo sapiens hypothetical protein FLJ23282 (FLJ23282), mRNA
NM 024803	Homo sapiens hypothetical protein FLJ21665 (FLJ21665), mRNA
NM_024795	Homo sapiens hypothetical protein FLJ22800 (FLJ22800), mRNA
NM 024767	Homo sapiens hypothetical protein FLJ21120 (FLJ21120), mRNA
NM 024760	Homo sapiens hypothetical protein FLJ14009 (FLJ14009), mRNA
NM 024741	Homo sapiens hypothetical protein FLJ12827 (FLJ12827), mRNA
NM 024723	Homo sapiens hypothetical protein FLJ23471 (FLJ23471), mRNA
NM 024720	Homo sapiens hypothetical protein FLJ23510 (FLJ23510), mRNA
NM 024698	Homo sapiens hypothetical protein FLJ13044 (FLJ13044), mRNA
NM_024692	Homo sapiens hypothetical protein FLJ21069 (FLJ21069), mRNA
NM_024689	Homo sapiens hypothetical protein FLJ14103 (FLJ14103), mRNA
NM_024687	Homo sapiens hypothetical protein FLJ23049 (FLJ23049), mRNA
NM_024648	Homo sapiens hypothetical protein FLJ222222 (FLJ222222), mRNA
NM_024622	Homo sapiens hypothetical protein FLJ21901 (FLJ21901), mRNA
NM_024611	Homo sapiens hypothetical protein FLJ11896 (FLJ11896), mRNA
NM_024591	Homo sapiens hypothetical protein FLJ11749 (FLJ11749), mRNA
NM_024561	Homo sapiens hypothetical protein FLJ22054 (FLJ22054), mRNA
NM_024540	Homo sapiens hypothetical protein FLJ20917 (FLJ20917), mRNA
NM_024518	Homo sapiens UL16-binding protein 3 (ULBP3), mRNA
NM_024515	Homo sapiens hypothetical protein MGC4645 (MGC4645), mRNA
NM_024504	Homo sapiens PR domain containing 14 (PRDM14), mRNA
NM_024501	Homo sapiens homeo box D1 (HOXD1), mRNA
NM_006821	Homo sapiens peroxisomal long-chain acyl-coA thioesterase (ZAP128), mRNA
NM_006680	Homo sapiens malic enzyme 3, NADP(+)-dependent, mitochondrial (ME3),
	mRNA
NM_001944	Homo sapiens desmoglein 3 (pemphigus vulgaris antigen) (DSG3), mRNA
NM_001943	Homo sapiens desmoglein 2 (DSG2), mRNA
NM_001942	Homo sapiens desmoglein 1 (DSG1), mRNA
NM_024500	Homo sapiens likely ortholog of mouse polydom (POLYDOM), mRNA
NM_024498	Homo sapiens zinc finger protein 117 (HPF9) (ZNF117), mRNA
NM_018943	Homo sapiens tubulin, alpha-like 2 (TUBAL2), mRNA
NM_015640	Homo sapiens PAI-1 mRNA-binding protein (PAI-RBP1), mRNA

NM_015332	Homo sapiens KIAA1068 protein (KIAA1068), mRNA
NM_022001	Homo sapiens SMAD in the antisense orientation (DAMS), mRNA
NM_021708	Homo sapiens leukocyte-associated Ig-like receptor 1 (LAIR1), transcript variant d, mRNA
NM_021706	Homo sapiens leukocyte-associated Ig-like receptor 1 (LAIR1), transcript variant b, mRNA
NM_002287	Homo sapiens leukocyte-associated Ig-like receptor 1 (LAIR1), transcript variant a, mRNA
NM 004424	Homo sapiens E4F transcription factor 1 (E4F1), mRNA
NM 018834	Homo sapiens matrin 3 (MATR3), mRNA
NM_017830	Homo sapiens ovarian carcinoma immunoreactive antigen (OCIA), mRNA
NM_006926	Homo sapiens surfactant, pulmonary-associated protein A2 (SFTPA2), mRNA
NM_005411	Homo sapiens surfactant, pulmonary-associated protein A1 (SFTPA1), mRNA
NM_024492	Homo sapiens apolipoprotein (a) related gene C (APOARGC), mRNA
NM_024491	Homo sapiens p10-binding protein (BITE), mRNA
NM_015472	Homo sapiens transcriptional co-activator with PDZ-binding motif (TAZ) (TAZ), mRNA
NM_017797	Homo sapiens BTB (POZ) domain containing 2 (BTBD2), mRNA
NM_002826	Homo sapiens quiescin Q6 (QSCN6), mRNA
NM_024010	Homo sapiens 5-methyltetrahydrofolate-homocysteine methyltransferase
	reductase (MTRR), transcript variant 2, mRNA
NM_004972	Homo sapiens Janus kinase 2 (a protein tyrosine kinase) (JAK2), mRNA
NM_000761	Homo sapiens cytochrome P450, subfamily I (aromatic compound-inducible), polypeptide 2 (CYP1A2), mRNA
NM_000104	Homo sapiens cytochrome P450, subfamily I (dioxin-inducible), polypeptide 1 (glaucoma 3, primary infantile) (CYP1B1), mRNA
NM_000499	Homo sapiens cytochrome P450, subfamily I (aromatic compound-inducible), polypeptide 1 (CYP1A1), mRNA
NM_024318	Homo sapiens immunoglobulin-like transcript 8 (ILT8), mRNA
NM_021806	Homo sapiens 2.19 gene (2.19), mRNA
NM_006208	Homo sapiens ectonucleotide pyrophosphatase/phosphodiesterase 1 (ENPP1), mRNA
NM_007076	Homo sapiens Huntingtin interacting protein E (HYPE), mRNA
NM_018571	Homo sapiens amyotrophic lateral sclerosis 2 (juvenile) chromosome region, candidate 2 (ALS2CR2), mRNA
NM_015049	Homo sapiens amyotrophic lateral sclerosis 2 (juvenile) chromosome region, candidate 3 (ALS2CR3), mRNA
NM_023036	Homo sapiens dynein intermediate chain 2 (DNAI2), mRNA
NM 022171	Homo sapiens T-cell leukemia translocation altered gene (TCTA), mRNA
NM 016128	Homo sapiens coat protein gamma-cop (LOC51137), mRNA
NM_021999	Homo sapiens integral membrane protein 2B (ITM2B), mRNA
NM_021992	Homo sapiens thymosin, beta, identified in neuroblastoma cells (TMSNB), mRNA
NM_021994	Homo sapiens zinc finger protein 277 (ZNF277), mRNA
NM_007257	Homo sapiens paraneoplastic antigen MA2 (PNMA2), mRNA
NM_021972	Homo sapiens sphingosine kinase 1 (SPHK1), mRNA
NM_021976	Homo sapiens retinoid X receptor, beta (RXRB), mRNA
NM_021963	Homo sapiens nucleosome assembly protein 1-like 2 (NAP1L2), mRNA
NM_021978	Homo sapiens suppression of tumorigenicity 14 (colon carcinoma, matriptase, epithin) (ST14), mRNA
NM_021977	Homo sapiens solute carrier family 22 (extraneuronal monoamine transporter), member 3 (SLC22A3), mRNA

NM_021964	Homo sapiens zinc finger protein 148 (pHZ-52) (ZNF148), mRNA
NM_021966	Homo sapiens T-cell leukemia/lymphoma 1A (TCL1A), mRNA
NM_012186	Homo sapiens forkhead box E3 (FOXE3), mRNA
NM_012182	Homo sapiens forkhead box B1 (FOXB1), mRNA
NM_006893	Homo sapiens ligatin (LGTN), mRNA
NM_021955	Homo sapiens guanine nucleotide binding protein (G protein), gamma
_	transducing activity polypeptide 1 (GNGT1), mRNA
NM_021959	Homo sapiens protein phosphatase 1, regulatory (inhibitor) subunit 11
	(PPP1R11), mRNA
NM_021951	Homo sapiens doublesex and mab-3 related transcription factor 1 (DMRT1),
	mRNA
NM_021960	Homo sapiens myeloid cell leukemia sequence 1 (BCL2-related) (MCL1),
	mRNA
NM_021952	Homo sapiens ELAV (embryonic lethal, abnormal vision, Drosophila)-like 4 (Hu
	antigen D) (ELAVL4), mRNA
NM_021949	Homo sapiens ATPase, Ca++ transporting, plasma membrane 3 (ATP2B3),
	mRNA
NM_021953	Homo sapiens forkhead box M1 (FOXM1), mRNA
NM_021956	Homo sapiens glutamate receptor, ionotropic, kainate 2 (GRIK2), mRNA
NM_004886	Homo sapiens amyloid beta (A4) precursor protein-binding, family A, member 3
37.6.006557	(X11-like 2) (APBA3), mRNA
NM_006557	Homo sapiens doublesex and mab-3 related transcription factor 2 (DMRT2),
) D. C. 000050	mRNA
NM_002253	Homo sapiens kinase insert domain receptor (a type III receptor tyrosine kinase)
ND 4 000170	(KDR), mRNA
NM_002178	Homo sapiens insulin-like growth factor binding protein 6 (IGFBP6), mRNA Homo sapiens succinate-CoA ligase, ADP-forming, beta subunit (SUCLA2),
NM_003850	mRNA
NM 003802	Homo sapiens myosin, heavy polypeptide 13, skeletal muscle (MYH13), mRNA
NM 006958	Homo sapiens zinc finger protein 16 (KOX 9) (ZNF16), mRNA
NM 006852	Homo sapiens tousled-like kinase 2 (TLK2), mRNA
NM 021229	Homo sapiens netrin 4 (NTN4), mRNA
NM 015718	Homo sapiens NADPH oxidase 3 (NOX3), mRNA
NM 015003	Homo sapiens golgin-67 (KIAA0855), mRNA
NM 006178	Homo sapiens N-ethylmaleimide-sensitive factor (NSF), mRNA
NM 003116	Homo sapiens sperm associated antigen 4 (SPAG4), mRNA
NM 018724	Homo sapiens interleukin 20 (IL20), mRNA
NM 019083	Homo sapiens hypothetical protein (FLJ10287), mRNA
NM 003114	Homo sapiens sperm associated antigen 1 (SPAG1), mRNA
NM 021097	Homo sapiens solute carrier family 8 (sodium/calcium exchanger), member 1
,,	(SLC8A1), mRNA
NM_021102	Homo sapiens serine protease inhibitor, Kunitz type, 2 (SPINT2), mRNA
NM 021101	Homo sapiens claudin 1 (CLDN1), mRNA
NM 021095	Homo sapiens solute carrier family 5 (sodium-dependent vitamin transporter),
	member 6 (SLC5A6), mRNA
NM 021076	Homo sapiens neurofilament, heavy polypeptide (200kD) (NEFH), mRNA
NM 001751	Homo sapiens cysteinyl-tRNA synthetase (CARS), mRNA
NM 021074	Homo sapiens NADH dehydrogenase (ubiquinone) flavoprotein 2 (24kD)
	(NDUFV2), mRNA
NM 020998	Homo sapiens macrophage stimulating 1 (hepatocyte growth factor-like)
1	(MST1), mRNA
NM_003147	Homo sapiens synovial sarcoma, X breakpoint 2 (SSX2), mRNA

NM_021964	Homo sapiens zinc finger protein 148 (pHZ-52) (ZNF148), mRNA
NM_021966	Homo sapiens T-cell leukemia/lymphoma 1A (TCL1A), mRNA
NM_012186	Homo sapiens forkhead box E3 (FOXE3), mRNA
NM_012182	Homo sapiens forkhead box B1 (FOXB1), mRNA
NM_006893	Homo sapiens ligatin (LGTN), mRNA
NM_021955	Homo sapiens guanine nucleotide binding protein (G protein), gamma
	transducing activity polypeptide 1 (GNGT1), mRNA
NM_021959	Homo sapiens protein phosphatase 1, regulatory (inhibitor) subunit 11
	(PPP1R11), mRNA
NM_021951	Homo sapiens doublesex and mab-3 related transcription factor 1 (DMRT1),
	mRNA
NM_021960	Homo sapiens myeloid cell leukemia sequence 1 (BCL2-related) (MCL1),
	mRNA
NM_021952	Homo sapiens ELAV (embryonic lethal, abnormal vision, Drosophila)-like 4 (Hu
	antigen D) (ELAVL4), mRNA
NM_021949	Homo sapiens ATPase, Ca++ transporting, plasma membrane 3 (ATP2B3),
	mRNA
NM_021953	Homo sapiens forkhead box M1 (FOXM1), mRNA
NM_021956	Homo sapiens glutamate receptor, ionotropic, kainate 2 (GRIK2), mRNA
NM_004886	Homo sapiens amyloid beta (A4) precursor protein-binding, family A, member 3
	(X11-like 2) (APBA3), mRNA
NM_006557	Homo sapiens doublesex and mab-3 related transcription factor 2 (DMRT2),
	mRNA
NM_002253	Homo sapiens kinase insert domain receptor (a type III receptor tyrosine kinase)
	(KDR), mRNA
NM_002178	Homo sapiens insulin-like growth factor binding protein 6 (IGFBP6), mRNA
NM_003850	Homo sapiens succinate-CoA ligase, ADP-forming, beta subunit (SUCLA2),
	mRNA
NM_003802	Homo sapiens myosin, heavy polypeptide 13, skeletal muscle (MYH13), mRNA
NM_006958	Homo sapiens zinc finger protein 16 (KOX 9) (ZNF16), mRNA
NM_006852	Homo sapiens tousled-like kinase 2 (TLK2), mRNA
NM_021229	Homo sapiens netrin 4 (NTN4), mRNA
NM_015718	Homo sapiens NADPH oxidase 3 (NOX3), mRNA
NM_015003	Homo sapiens golgin-67 (KIAA0855), mRNA
NM_006178	Homo sapiens N-ethylmaleimide-sensitive factor (NSF), mRNA
NM_003116	Homo sapiens sperm associated antigen 4 (SPAG4), mRNA
NM_018724	Homo sapiens interleukin 20 (IL20), mRNA
NM_019083	Homo sapiens hypothetical protein (FLJ10287), mRNA
NM_003114	Homo sapiens sperm associated antigen 1 (SPAG1), mRNA
NM_021097	Homo sapiens solute carrier family 8 (sodium/calcium exchanger), member 1
	(SLC8A1), mRNA
NM_021102	Homo sapiens serine protease inhibitor, Kunitz type, 2 (SPINT2), mRNA
NM_021101	Homo sapiens claudin 1 (CLDN1), mRNA
NM_021095	Homo sapiens solute carrier family 5 (sodium-dependent vitamin transporter),
	member 6 (SLC5A6), mRNA
NM_021076	Homo sapiens neurofilament, heavy polypeptide (200kD) (NEFH), mRNA
NM_001751	Homo sapiens cysteinyl-tRNA synthetase (CARS), mRNA
NM_021074	Homo sapiens NADH dehydrogenase (ubiquinone) flavoprotein 2 (24kD)
	(NDUFV2), mRNA
NM_020998	Homo sapiens macrophage stimulating 1 (hepatocyte growth factor-like)
	(MST1), mRNA
NM 003147	Homo sapiens synovial sarcoma, X breakpoint 2 (SSX2), mRNA

NM_015392	Homo sapiens neural proliferation, differentiation and control, 1 (NPDC1), mRNA
NM 020482	Homo sapiens activator of CREM in testis (ACT), mRNA
NM 014509	Homo sapiens kraken-like (BK126B4.1), mRNA
NM_005132	Homo sapiens Rec8p, a meiotic recombination and sister chromatid cohesion phosphoprotein of the rad21p family (REC8), mRNA
NM 018896	Homo sapiens calcium channel, voltage-dependent, alpha 1G subunit
	(CACNA1G), mRNA
NM_005329	Homo sapiens hyaluronan synthase 3 (HAS3), mRNA
NM_015193	Homo sapiens activity-regulated cytoskeleton-associated protein (ARC), mRNA
NM_016203	Homo sapiens protein kinase, AMP-activated, gamma 2 non-catalytic subunit (PRKAG2), mRNA
NM_000627	Homo sapiens latent transforming growth factor beta binding protein 1 (LTBP1),
ND4 002454	mRNA Homo sapiens 5-methyltetrahydrofolate-homocysteine methyltransferase
NM_002454	reductase (MTRR), transcript variant 1, mRNA
NM_001091	Homo sapiens amiloride binding protein 1 (amine oxidase (copper-containing))
	(ABP1), mRNA
NM_024016	Homo sapiens homeo box B8 (HOXB8), mRNA
NM_024015	Homo sapiens homeo box B4 (HOXB4), mRNA
NM_015227	Homo sapiens KIAA0958 protein (KIAA0958), mRNA
NM_024430	Homo sapiens proline-serine-threonine phosphatase interacting protein 2
	(PSTPIP2), mRNA
NM_003588	Homo sapiens cullin 4B (CUL4B), mRNA
NM_016059	Homo sapiens peptidylprolyl isomerase (cyclophilin)-like 1 (PPIL1), mRNA
NM_014432	Homo sapiens interleukin 20 receptor, alpha (IL20RA), mRNA
NM_000270	Homo sapiens nucleoside phosphorylase (NP), mRNA
NM_003021	Homo sapiens small glutamine-rich tetratricopeptide repeat (TPR)-containing (SGT), mRNA
NM_002038	Homo sapiens interferon, alpha-inducible protein (clone IFI-6-16) (G1P3),
NIM 022972	transcript variant 1, mRNA
NM_022873	Homo sapiens interferon, alpha-inducible protein (clone IFI-6-16) (G1P3), transcript variant 3, mRNA
NM_022872	Homo sapiens interferon, alpha-inducible protein (clone IFI-6-16) (G1P3), transcript variant 2, mRNA
NM_022803	Homo sapiens uncoupling protein 3 (mitochondrial, proton carrier) (UCP3),
	transcript variant short, nuclear gene encoding mitochondrial protein, mRNA
NM 003356	Homo sapiens uncoupling protein 3 (mitochondrial, proton carrier) (UCP3),
	transcript variant long, nuclear gene encoding mitochondrial protein, mRNA
NM_022810	Homo sapiens solute carrier family 25 (mitochondrial carrier, brain), member 14
	(SLC25A14), transcript variant short, nuclear gene encoding mitochondrial
	protein, mRNA
NM 003355	Homo sapiens uncoupling protein 2 (mitochondrial, proton carrier) (UCP2),
	nuclear gene encoding mitochondrial protein, mRNA
NM_021833	Homo sapiens uncoupling protein 1 (mitochondrial, proton carrier) (UCP1),
	nuclear gene encoding mitochondrial protein, mRNA
NM_002231	Homo sapiens kangai 1 (suppression of tumorigenicity 6, prostate; CD82 antigen
	(R2 leukocyte antigen, antigen detected by monoclonal and antibody IA4))
	(KAII), mRNA
NM_004967	Homo sapiens integrin-binding sialoprotein (bone sialoprotein, bone sialoprotein II) (IBSP), mRNA
NM 000490	Homo sapiens arginine vasopressin (neurophysin II, antidiuretic hormone,
	and any time and the second discussion in minimum in minimum,

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ND 6 000000	diabetes insipidus, neurohypophyseal) (AVP), mRNA Homo sapiens survival of motor neuron 2, centromeric (SMN2), transcript
NM_022877	
27.6.000000	variant c, mRNA Homo sapiens survival of motor neuron 2, centromeric (SMN2), transcript
NM_022876	
37 6 000075	variant b, mRNA Homo sapiens survival of motor neuron 2, centromeric (SMN2), transcript
NM_022875	
27 6 017411	variant a, mRNA Homo sapiens survival of motor neuron 2, centromeric (SMN2), transcript
NM_017411	variant d, mRNA
NM 005474	Homo sapiens histone deacetylase 5 (HDAC5), mRNA
NM 006037	Homo sapiens histone deacetylase 4 (HDAC4), mRNA
	Homo sapiens a disintegrin and metalloproteinase domain 12 (meltrin alpha)
NM_003474	(ADAM12), transcript variant 1, mRNA
NM 000344	Homo sapiens survival of motor neuron 1, telomeric (SMN1), transcript variant
14141_000344	d, mRNA
NM 022874	Homo sapiens survival of motor neuron 1, telomeric (SMN1), transcript variant
14141_022074	b, mRNA
NM 006400	Homo sapiens dynactin 2 (p50) (DCTN2), mRNA
NM 021969	Homo sapiens nuclear receptor subfamily 0, group B, member 2 (NR0B2),
	mRNA
NM 021967	Homo sapiens small EDRK-rich factor 1A (telomeric) (SERF1A), mRNA
NM 001515	Homo sapiens general transcription factor IIH, polypeptide 2 (44kD subunit)
1111_001010	(GTF2H2), mRNA
NM 003951	Homo sapiens solute carrier family 25 (mitochondrial carrier, brain), member 14
	(SLC25A14), transcript variant long, nuclear gene encoding mitochondrial
	protein, mRNA
NM 004277	Homo sapiens uncoupling protein 4 (UCP4), nuclear gene encoding
_	mitochondrial protein, mRNA
NM_004536	Homo sapiens baculoviral IAP repeat-containing 1 (BIRC1), mRNA
NM_000346	Homo sapiens SRY (sex determining region Y)-box 9 (campomelic dysplasia,
	autosomal sex-reversal) (SOX9), mRNA
NM_003645	Homo sapiens fatty-acid-Coenzyme A ligase, very long-chain 1 (FACVL1),
	mRNA
NM_024409	Homo sapiens natriuretic peptide precursor C (NPPC), mRNA
NM_024410	Homo sapiens outer dense fibre of sperm tails 1 (ODF1), mRNA
NM_004180	Homo sapiens TRAF family member-associated NFKB activator (TANK),
***	mRNA
NM_024332	Homo sapiens c6.1A (C6.1A), mRNA
NM_024324	Homo sapiens hypothetical protein MGC11256 (MGC11256), mRNA
NM_024315	Homo sapiens hypothetical protein MGC4175 (MGC4175), mRNA
NM_024311	Homo sapiens hypothetical protein ET (ET), mRNA
NM_024309	Homo sapiens hypothetical protein MGC4289 (MGC4289), mRNA
NM_024306	Homo sapiens fatty acid hydroxylase (FAAH), mRNA
NM_024300	Homo sapiens hypothetical protein MGC2217 (MGC2217), mRNA
NM_024296	Homo sapiens hypothetical protein MGC1203 (MGC1203), mRNA
NM_024294	Homo sapiens hypothetical protein MGC4614 (MGC4614), mRNA
NM_024292	Homo sapiens ubiquitin-like 5 (UBL5), mRNA
NM_024012	Homo sapiens 5-hydroxytryptamine (serotonin) receptor 5A (HTR5A), mRNA
NM_024123	Homo sapiens putative Ly-6 superfamily member (G6E), mRNA
NM_021904	Homo sapiens gamma-aminobutyric acid (GABA) B receptor, 1 (GABBR1),
	transcript variant 3, mRNA
NM_021903	Homo sapiens gamma-aminobutyric acid (GABA) B receptor, 1 (GABBR1),

NB4 001470	transcript variant 2, mRNA
NM_001470	Homo sapiens gamma-aminobutyric acid (GABA) B receptor, 1 (GABBR1), transcript variant 1, mRNA
NM_001858	Homo sapiens collagen, type XIX, alpha 1 (COL19A1), mRNA
NM_015071	Homo sapiens GTPase regulator associated with the focal adhesion kinase
	pp125(FAK); KIAA0621 protein (KIAA0621), mRNA
NM_007329	Homo sapiens deleted in malignant brain tumors 1 (DMBT1), transcript variant 2, mRNA
NM_023004	Homo sapiens nogo receptor (NOGOR), mRNA
NM_005371	Homo sapiens methyltransferase-like 1 (METTL1), transcript variant 1, mRNA
NM_023033	Homo sapiens methyltransferase-like 1 (METTL1), transcript variant 3, mRNA
NM_023032	Homo sapiens methyltransferase-like 1 (METTL1), transcript variant 2, mRNA
NM_014289	Homo sapiens calpain 6 (CAPN6), mRNA
NM_023089	Homo sapiens calpain 10 (CAPN10), transcript variant 7, mRNA
NM_023088	Homo sapiens calpain 10 (CAPN10), transcript variant 6, mRNA
NM_023087	Homo sapiens calpain 10 (CAPN10), transcript variant 5, mRNA
NM_023086	Homo sapiens calpain 10 (CAPN10), transcript variant 4, mRNA
NM_023085	Homo sapiens calpain 10 (CAPN10), transcript variant 3, mRNA
NM_023084	Homo sapiens calpain 10 (CAPN10), transcript variant 2, mRNA
NM_023083	Homo sapiens calpain 10 (CAPN10), transcript variant 1, mRNA
NM_021251	Homo sapiens calpain 10 (CAPN10), transcript variant 8, mRNA
NM_005083	Homo sapiens U2 small nuclear ribonucleoprotein auxiliary factor, small subunit 1 (U2AF1RS1), mRNA
NM_023031	Homo sapiens fibroblast growth factor receptor 2 (bacteria-expressed kinase,
_	keratinocyte growth factor receptor, craniofacial dysostosis 1. Crouzon
•	syndrome, Pfeiffer syndrome, Jackson-Weiss syndrome) (FGFR2), transcript
375 0770	variant 13, mRNA
NM_023030	Homo sapiens fibroblast growth factor receptor 2 (bacteria-expressed kinase,
	keratinocyte growth factor receptor, craniofacial dysostosis 1 Crouzon
	syndrome, Pfeiffer syndrome, Jackson-Weiss syndrome) (FGFR2), transcript variant 12, mRNA
NM_023028	Homo sapiens fibroblast growth factor receptor 2 (bacteria-expressed kinase,
**	keratinocyte growth factor receptor, craniofacial dysostosis 1. Croppon
	syndrome, Pfeiffer syndrome, Jackson-Weiss syndrome) (FGFR2) transcript
	variant 10, mr.NA
NM_022976	Homo sapiens fibroblast growth factor receptor 2 (bacteria-expressed kinase,
	Reratinocyte growth factor receptor, craniofacial dysostosis 1 Crouzon
	syndrome, Pteriffer syndrome, Jackson-Weiss syndrome) (FGFR2), transcript
NM 022075	variant 9, mich A
NM_022975	Homo sapiens fibroblast growth factor receptor 2 (bacteria-expressed kinase,
	keratinocyte growth factor receptor, craniofacial dysostosis 1, Crouzon
	syndrome, Pfeiffer syndrome, Jackson-Weiss syndrome) (FGFR2), transcript variant 8, mRNA
NM_022974	
1111_022774	Homo sapiens fibroblast growth factor receptor 2 (bacteria-expressed kinase,
	keratinocyte growth factor receptor, craniofacial dysostosis 1, Crouzon
	syndrome, Pfeiffer syndrome, Jackson-Weiss syndrome) (FGFR2), transcript variant 7, mRNA
NM_022973	
	Homo sapiens fibroblast growth factor receptor 2 (bacteria-expressed kinase,
ļ	keratinocyte growth factor receptor, craniofacial dysostosis 1, Crouzon syndrome, Pfeiffer syndrome, Jackson-Weiss syndrome) (FGFR2), transcript
	variant 6, mRNA
NM_022972	Homo sapiens fibroblast growth factor receptor 2 (bacteria-expressed kinase,
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	keratinocyte growth factor receptor, craniofacial dysostosis 1, Crouzon
	syndrome, Pfeiffer syndrome, Jackson-Weiss syndrome) (FGFR2), transcript variant 5, mRNA
NM_022971	Homo sapiens fibroblast growth factor receptor 2 (bacteria-expressed kinase,
-	keratinocyte growth factor receptor, craniofacial dysostosis 1, Crouzon
	syndrome, Pfeiffer syndrome, Jackson-Weiss syndrome) (FGFR2), transcript
	variant 4, mRNA
NM_022970	Homo sapiens fibroblast growth factor receptor 2 (bacteria-expressed kinase,
	keratinocyte growth factor receptor, craniofacial dysostosis 1, Crouzon
	syndrome, Pfeiffer syndrome, Jackson-Weiss syndrome) (FGFR2), transcript
\\\\\\\\\\\\\\\\\\\\\\\\\\\\\\\\\\\\\\	variant 3, mRNA
NM_022969	Homo sapiens fibroblast growth factor receptor 2 (bacteria-expressed kinase,
	keratinocyte growth factor receptor, craniofacial dysostosis 1, Crouzon
	syndrome, Pfeiffer syndrome, Jackson-Weiss syndrome) (FGFR2), transcript variant 2, mRNA
NM_015850	Homo sapiens fibroblast growth factor receptor 1 (fms-related tyrosine kinase 2,
	Pfeiffer syndrome) (FGFR1), transcript variant 2, mRNA
NM_023111	Homo sapiens fibroblast growth factor receptor 1 (fms-related tyrosine kinase 2,
	Pfeiffer syndrome) (FGFR1), transcript variant 9, mRNA
NM_023110	Homo sapiens fibroblast growth factor receptor 1 (fms-related tyrosine kinase 2,
>D 6 000100	Pfeiffer syndrome) (FGFR1), transcript variant 8, mRNA
NM_023109	Homo sapiens fibroblast growth factor receptor 1 (fms-related tyrosine kinase 2,
ND (022020	Pfeiffer syndrome) (FGFR1), transcript variant 7, mRNA
NM_023029	Homo sapiens fibroblast growth factor receptor 2 (bacteria-expressed kinase,
	keratinocyte growth factor receptor, craniofacial dysostosis 1, Crouzon
	syndrome, Pfeiffer syndrome, Jackson-Weiss syndrome) (FGFR2), transcript variant 11, mRNA
NM_023108	Homo sapiens fibroblast growth factor receptor 1 (fms-related tyrosine kinase 2,
	Pfeiffer syndrome) (FGFR1), transcript variant 6, mRNA
NM 000141	Homo sapiens fibroblast growth factor receptor 2 (bacteria-expressed kinase,
_	keratinocyte growth factor receptor, craniofacial dysostosis 1, Crouzon
	syndrome, Pfeiffer syndrome, Jackson-Weiss syndrome) (FGFR2), transcript
	variant 1, mRNA
NM_023107	Homo sapiens fibroblast growth factor receptor 1 (fms-related tyrosine kinase 2,
	Pfeiffer syndrome) (FGFR1), transcript variant 5, mRNA
NM_023106	Homo sapiens fibroblast growth factor receptor 1 (fms-related tyrosine kinase 2,
ND (000105	Pfeiffer syndrome) (FGFR1), transcript variant 4, mRNA
NM_023105	Homo sapiens fibroblast growth factor receptor 1 (fms-related tyrosine kinase 2,
ND4 000604	Pfeiffer syndrome) (FGFR1), transcript variant 3, mRNA
NM_000604	Homo sapiens fibroblast growth factor receptor 1 (fms-related tyrosine kinase 2,
NM 024018	Pfeiffer syndrome) (FGFR1), transcript variant 1, mRNA
NM 017614	Homo sapiens butyrophilin, subfamily 2, member A3 (BTN2A3), mRNA
NM 005434	Homo sapiens betaine-homocysteine methyltransferase 2 (BHMT2), mRNA Homo sapiens BENE protein (BENE), mRNA
NM 000351	Homo sapiens steroid sulfatase (microsomal), arylsulfatase C, isozyme S (STS),
14142_000551	mRNA
NM_024105	Homo sapiens hypothetical protein MGC3136 (MGC3136), mRNA
NM_024098	Homo sapiens hypothetical protein MGC2574 (MGC2574), mRNA
NM_024096	Homo sapiens hypothetical protein MGC5627 (MGC5627), mRNA
NM_024095	Homo sapiens hypothetical protein MGC5540 (MGC5540), mRNA
NM_024091	Homo sapiens hypothetical protein MGC5297 (MGC5297), mRNA
NM 024089	

NM_024082	Homo sapiens transmembrane gamma-carboxyglutamic acid protein 3 (TMG3), mRNA
NM_024081	Homo sapiens transmembrane gamma-carboxyglutamic acid protein 4 (TMG4), mRNA
NM_024079	Homo sapiens hypothetical protein MGC2840 similar to a putative glucosyltransferase (MGC2840), mRNA
NM 024078	Homo sapiens hypothetical protein MGC3162 (MGC3162), mRNA
NM 024075	Homo sapiens LENG5 protein (LENG5), mRNA
NM 024073	Homo sapiens hypothetical protein MGC2875 (MGC2875), mRNA
NM 024060	Homo sapiens hypothetical protein MGC2873 (MGC2873), mRNA
NM 024056	Homo sapiens hypothetical protein MGC5576 (MGC5576), mRNA
NM 024054	Homo sapiens hypothetical protein MGC3576 (MGC3576), mRNA
NM 024051	Homo sapiens hypothetical protein MGC3077 (MGC3077), mRNA
NM 024047	Homo sapiens hypothetical protein MGC3037 (MGC3037), mRNA
NM 024044	Homo sapiens hypothetical protein MGC5178 (MGC5037), mRNA
NM_024043	Homo sapiens hypothetical protein MGC3101 (MGC3101), mRNA
NM 024035	Homo sapiens hypothetical protein MGC3113 (MGC3113), mRNA
NM_024034	Homo sapiens hypothetical protein MGC3129 similar to ganglioside-induced
-	differentiation-associated protein (MGC3129), mRNA
NM_024009	Homo sapiens gap junction protein, beta 3, 31kD (connexin 31) (GJB3), mRNA
NM_024013	Homo sapiens interferon, alpha 1 (IFNA1), mRNA
NM_000521	Homo sapiens hexosaminidase B (beta polypeptide) (HEXB), mRNA
NM_000520	Homo sapiens hexosaminidase A (alpha polypeptide) (HEXA), mRNA
NM_006044	Homo sapiens histone deacetylase 6 (HDAC6), mRNA
NM_003883	Homo sapiens histone deacetylase 3 (HDAC3), mRNA
NM_004964	Homo sapiens histone deacetylase 1 (HDAC1), mRNA
NM_001492	Homo sapiens growth differentiation factor 1 (GDF1), mRNA
NM_018486	Homo sapiens histone deacetylase 8 (HDAC8), mRNA
NM_005089	Homo sapiens U2 small nuclear ribonucleoprotein auxiliary factor, small subunit 2 (U2AF1RS2), mRNA
NM_004285	Homo sapiens hexose-6-phosphate dehydrogenase (glucose 1-dehydrogenase) (H6PD), mRNA
NM_007210	Homo sapiens UDP-N-acetyl-alpha-D-galactosamine:polypeptide N-
	acetylgalactosaminyltransferase 6 (GalNAc-T6) (GALNT6), mRNA
NM_003774	Homo sapiens UDP-N-acetyl-alpha-D-galactosamine:polyneptide N-
	acetylgalactosaminyltransferase 4 (GalNAc-T4) (GALNT4), mRNA
NM_020474	Homo sapiens UDP-N-acetyl-alpha-D-galactosamine:polypeptide N-
	acetylgalactosaminyltransferase 1 (GalNAc-T1) (GALNT1), mRNA
NM_015507	Homo sapiens EGF-like-domain, multiple 6 (EGFL6), mRNA
NM_004942	Homo sapiens defensin, beta 2 (DEFB2), mRNA
NM_005218	Homo sapiens defensin, beta 1 (DEFB1), mRNA
NM_002474	Homo sapiens myosin, heavy polypeptide 11, smooth muscle (MYH11),
) T (000000	transcript variant SM1, mRNA
NM_022870	Homo sapiens myosin, heavy polypeptide 11, smooth muscle (MYH11),
ND 6 000044	transcript variant SM3, mRNA
NM_022844	Homo sapiens myosin, heavy polypeptide 11, smooth muscle (MYH11), transcript variant SM2, mRNA
NM_001755	Homo sapiens core-binding factor, beta subunit (CBFB), transcript variant 2,
	mRNA
NM_016458	Homo sapiens hypothetical protein (LOC51236), mRNA
NM_020836	Homo sapiens KIAA1446 protein (KIAA1446), mRNA
NM_015407	Homo sapiens DKFZP564O243 protein (DKFZP564O243), mRNA
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NM_015062	Homo sapiens KIAA0595 protein (KIAA0595), mRNA
NM_019100	Homo sapiens DNA methyltransferase 1-associated protein 1 (DMAP1), mRNA
NM_015442	Homo sapiens hypothetical protein FLJ12890 (FLJ12890), mRNA
NM_023948	Homo sapiens hypothetical protein AF053356_CDS3 (AF053356_CDS3), mRNA
NM_022036	Homo sapiens G protein-coupled receptor, family C, group 5, member C (GPRC5C), transcript variant 1, mRNA
NM_018653	Homo sapiens G protein-coupled receptor, family C, group 5, member C (GPRC5C), transcript variant 2, mRNA
NM 000707	Homo sapiens arginine vasopressin receptor 1B (AVPR1B), mRNA
NM 000706	Homo sapiens arginine vasopressin receptor 1A (AVPR1A), mRNA
NM 021923	Homo sapiens fibroblast growth factor receptor-like 1 (FGFRL1), mRNA
NM_002011	Homo sapiens fibroblast growth factor receptor 4 (FGFR4), transcript variant 1, mRNA
NM_022963	Homo sapiens fibroblast growth factor receptor 4 (FGFR4), transcript variant 2, mRNA
NM_022965	Homo sapiens fibroblast growth factor receptor 3 (achondroplasia, thanatophoric dwarfism) (FGFR3), transcript variant 2, mRNA
NM_000142	Homo sapiens fibroblast growth factor receptor 3 (achondroplasia, thanatophoric dwarfism) (FGFR3), transcript variant 1, mRNA
NM_022336	Homo sapiens ectodysplasin 1, anhidrotic receptor (EDAR), mRNA
NM_018654	Homo sapiens G protein-coupled receptor, family C, group 5, member D (GPRC5D), mRNA
NM_002534	Homo sapiens 2',5'-oligoadenylate synthetase 1 (40-46 kD) (OAS1), transcript variant E16, mRNA
NM_016816	Homo sapiens 2',5'-oligoadenylate synthetase 1 (40-46 kD) (OAS1), transcript variant E18, mRNA
NM 014501	Homo sapiens ubiquitin carrier protein (E2-EPF), mRNA
NM 000595	Homo sapiens lymphotoxin alpha (TNF superfamily, member 1) (LTA), mRNA
NM 007040	Homo sapiens E1B-55kDa-associated protein 5 (E1B-AP5), mRNA
NM 001232	Homo sapiens calsequestrin 2 (cardiac muscle) (CASQ2), mRNA
NM_001231	Homo sapiens calsequestrin 1 (fast-twitch, skeletal muscle) (CASQ1), nuclear gene encoding mitochondrial protein, mRNA
NM_003925	Homo sapiens methyl-CpG binding domain protein 4 (MBD4), mRNA
NM_002059	Homo sapiens growth hormone 2 (GH2), transcript variant 1, mRNA
NM 022558	Homo sapiens growth hormone 2 (GH2), transcript variant 3, mRNA
NM_022557	Homo sapiens growth hormone 2 (GH2), transcript variant 2, mRNA
NM_022556	Homo sapiens growth hormone 2 (GH2), transcript variant 4, mRNA
NM_022562	Homo sapiens growth hormone 1 (GH1), transcript variant 5, mRNA
NM 022561	Homo sapiens growth hormone 1 (GH1), transcript variant 4, mRNA
NM 022560	Homo sapiens growth hormone 1 (GH1), transcript variant 3, mRNA
NM 022559	Homo sapiens growth hormone 1 (GH1), transcript variant 2, mRNA
NM 000515	Homo sapiens growth hormone 1 (GH1), transcript variant 1, mRNA
NM 021801	Homo sapiens matrix metalloproteinase 26 (MMP26), mRNA
NM_022718	Homo sapiens matrix metalloproteinase 25 (MMP25), transcript variant 2, mRNA
NM_022468	Homo sapiens matrix metalloproteinase 25 (MMP25), transcript variant 1, mRNA
NM_006690	Homo sapiens matrix metalloproteinase 24 (membrane-inserted) (MMP24), mRNA
NM_004771	Homo sapiens matrix metalloproteinase 20 (enamelysin) (MMP20), mRNA
NM 002423	Homo sapiens matrix metalloproteinase 7 (matrilysin, uterine) (MMP7), mRNA
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NM 002422	Homo sapiens matrix metalloproteinase 3 (stromelysin 1, progelatinase)
_	(MMP3), mRNA
NM_005941	Homo sapiens matrix metalloproteinase 16 (membrane-inserted) (MMP16), transcript variant 1, mRNA
NM_022564	Homo sapiens matrix metalloproteinase 16 (membrane-inserted) (MMP16), transcript variant 2, mRNA
NM_002421	Homo sapiens matrix metalloproteinase 1 (interstitial collagenase) (MMP1), mRNA
NM_004995	Homo sapiens matrix metalloproteinase 14 (membrane-inserted) (MMP14), mRNA
NM_002427	Homo sapiens matrix metalloproteinase 13 (collagenase 3) (MMP13), mRNA
NM_005940	Homo sapiens matrix metalloproteinase 11 (stromelysin 3) (MMP11), mRNA
NM_022792	Homo sapiens matrix metalloproteinase 19 (MMP19), transcript variant rasi-9, mRNA
NM_022791	Homo sapiens matrix metalloproteinase 19 (MMP19), transcript variant rasi-6, mRNA
NM_022790	Homo sapiens matrix metalloproteinase 19 (MMP19), transcript variant rasi-3, mRNA
NM_002429	Homo sapiens matrix metalloproteinase 19 (MMP19), transcript variant rasi-1, mRNA
NM_004530	Homo sapiens matrix metalloproteinase 2 (gelatinase A, 72kD gelatinase, 72kD type IV collagenase) (MMP2), mRNA
NM_004994	Homo sapiens matrix metalloproteinase 9 (gelatinase B, 92kD gelatinase, 92kD type IV collagenase) (MMP9), mRNA
NM_004142	Homo sapiens matrix metalloproteinase-like 1 (MMPL1), mRNA
NM_002424	Homo sapiens matrix metalloproteinase 8 (neutrophil collagenase) (MMP8), mRNA
NM_002428	Homo sapiens matrix metalloproteinase 15 (membrane-inserted) (MMP15), mRNA
NM_002426	Homo sapiens matrix metalloproteinase 12 (macrophage elastase) (MMP12), mRNA
NM_002425	Homo sapiens matrix metalloproteinase 10 (stromelysin 2) (MMP10), mRNA
NM_022804	Homo sapiens SNRPN upstream reading frame (SNURF), transcript variant 2, mRNA
NM_005678	Homo sapiens SNRPN upstream reading frame (SNURF), transcript variant 1, mRNA
NM_003097	Homo sapiens small nuclear ribonucleoprotein polypeptide N (SNRPN), transcript variant 1, mRNA
NM_022808	Homo sapiens small nuclear ribonucleoprotein polypeptide N (SNRPN), transcript variant 5, mRNA
NM_022807	Homo sapiens small nuclear ribonucleoprotein polypeptide N (SNRPN), transcript variant 4, mRNA
NM_022806	Homo sapiens small nuclear ribonucleoprotein polypeptide N (SNRPN), transcript variant 3, mRNA
NM_022805	Homo sapiens small nuclear ribonucleoprotein polypeptide N (SNRPN), transcript variant 2, mRNA
NM_022717	Homo sapiens U1-snRNP binding protein homolog (70kD) (U1SNRNPBP), transcript variant 2, mRNA
NM_006759	Homo sapiens UDP-glucose pyrophosphorylase 2 (UGP2), mRNA
NM_001400	Homo sapiens endothelial differentiation, sphingolipid G-protein-coupled receptor, 1 (EDG1), mRNA
NM_005586	Homo sapiens MyoD family inhibitor (MDFI), mRNA

NM_022978	Homo sapiens small EDRK-rich factor 1B (centromeric) (SERF1B), mRNA
NM_023947	Homo sapiens hypothetical protein MGC3234 (MGC3234), mRNA
NM_023942	Homo sapiens hypothetical protein MGC3036 (MGC3036), mRNA
NM_023933	Homo sapiens hypothetical protein MGC2494 (MGC2494), mRNA
NM_005471	Homo sapiens glucosamine-6-phosphate isomerase (GNPI), mRNA
NM_023925	Homo sapiens hypothetical protein FLJ22569 (FLJ22569), mRNA
NM_004076	Homo sapiens crystallin, beta B3 (CRYBB3), mRNA
NM_015717	Homo sapiens Langerhans cell specific c-type lectin (LANGERIN), mRNA
NM_012329	Homo sapiens monocyte to macrophage differentiation-associated (MMD), mRNA
NM_007020	Homo sapiens U1-snRNP binding protein homolog (70kD) (U1SNRNPBP), transcript variant 1, mRNA
NM_006465	Homo sapiens dead ringer (Drosophila)-like 2 (bright and dead ringer) (DRIL2), mRNA
NM_000015	Homo sapiens N-acetyltransferase 2 (arylamine N-acetyltransferase) (NAT2), mRNA
NM_000496	Homo sapiens crystallin, beta B2 (CRYBB2), mRNA
NM_001886	Homo sapiens crystallin, beta A4 (CRYBA4), mRNA
NM_023080	Homo sapiens hypothetical protein FLJ20989 (FLJ20989), mRNA
NM_023039	Homo sapiens ankyrin repeat, family A (RFXANK-like), 2 (ANKRA2), mRNA
NM_021905	Homo sapiens gamma-aminobutyric acid (GABA) B receptor, 1 (GABBR1), transcript variant 4, mRNA
NM_020554	Homo sapiens T-cell leukemia/lymphoma 6 (TCL6), transcript variant TCL6d1, mRNA
NM_020553	Homo sapiens T-cell leukemia/lymphoma 6 (TCL6), transcript variant TCL6c1, mRNA
NM_020552	Homo sapiens T-cell leukemia/lymphoma 6 (TCL6), transcript variant TCL6b1, mRNA
NM_020550	Homo sapiens T-cell leukemia/lymphoma 6 (TCL6), transcript variant TCL6a3, mRNA
NM_012468	Homo sapiens T-cell leukemia/lymphoma 6 (TCL6), transcript variant TCL6a1, mRNA
NM_014418	Homo sapiens T-cell leukemia/lymphoma 6 (TCL6), transcript variant TCL6a2, mRNA
NM_016730	Homo sapiens folate receptor 1 (adult) (FOLR1), transcript variant 3, mRNA
NM_016729	Homo sapiens folate receptor 1 (adult) (FOLR1), transcript variant 4, mRNA
NM_016725	Homo sapiens folate receptor 1 (adult) (FOLR1), transcript variant 1, mRNA
NM_016724	Homo sapiens folate receptor 1 (adult) (FOLR1), transcript variant 7, mRNA
NM_016025	Homo sapiens CGI-81 protein (DREV1), mRNA
NM_004406	Homo sapiens deleted in malignant brain tumors 1 (DMBT1), transcript variant 1, mRNA
NM_000197	Homo sapiens hydroxysteroid (17-beta) dehydrogenase 3 (HSD17B3), mRNA
NM_001220	Homo sapiens calcium/calmodulin-dependent protein kinase (CaM kinase) II
	beta (CAMK2B), mRNA
NM_019071	Homo sapiens inhibitor of growth family, member 3 (ING3), mRNA
NM_016731	Homo sapiens folate receptor 1 (adult) (FOLR1), transcript variant 8, mRNA
NM_023018	Homo sapiens hypothetical protein FLJ13052 (FLJ13052), mRNA
NM_023016	Homo sapiens hypothetical protein FLJ21870 (FLJ21870), mRNA
NM_022911	Homo sapiens solute carrier family 26, member 6 (SLC26A6), mRNA
NM_021071	Homo sapiens ADP-ribosyltransferase 4 (ART4), mRNA
NM_022113	Homo sapiens kinesin family member 13A (KIF13A), mRNA
NM_012449	Homo sapiens six transmembrane epithelial antigen of the prostate (STEAP),
	J. T. M. Product (O. L. M. I.),

	DATA
ND 6 016512	mRNA Homo sapiens MAK-related kinase (KIAA0936), mRNA
NM_016513	Homo sapiens MAK-related kinase (KIAA0930), mRNA Homo sapiens MAK-related kinase (KIAA0936), mRNA
NM_014920	Homo sapiens max-related kinase (Kraausso), hikkya Homo sapiens related to the N terminus of tre (RNTRE), mRNA
NM_014688	Homo sapiens MLL septin-like fusion (MSF), mRNA
NM_006640	Homo sapiens TRK-fused gene (TFG), mRNA
NM_006070	
NM_004809	Homo sapiens stomatin-like 1 (STOML1), mRNA
NM_000297	Homo sapiens polycystic kidney disease 2 (autosomal dominant) (PKD2), mRNA
NM_016307	Homo sapiens paired related homeobox protein (PRX2), mRNA
NM_003924	Homo sapiens paired mesoderm homeobox 2b (PMX2B), mRNA
NM_006902	Homo sapiens paired mesoderm homeo box 1 (PMX1), transcript variant pmx-1a, mRNA
NM_022716	Homo sapiens paired mesoderm homeo box 1 (PMX1), transcript variant pmx-1b, mRNA
NM_000916	Homo sapiens oxytocin receptor (OXTR), mRNA
NM_000915	Homo sapiens oxytocin, prepro- (neurophysin I) (OXT), mRNA
NM_006188	Homo sapiens oncomodulin (OCM), mRNA
NM_022664	Homo sapiens extracellular matrix protein 1 (ECM1), transcript variant 2, mRNA
NM 004092	Homo sapiens enoyl Coenzyme A hydratase, short chain, 1, mitochondrial
_	(ECHS1), nuclear gene encoding mitochondrial protein, mRNA
NM_022652	Homo sapiens dual specificity phosphatase 6 (DUSP6), transcript variant 2, mRNA
NM 004419	Homo sapiens dual specificity phosphatase 5 (DUSP5), mRNA
NM_004425	Homo sapiens extracellular matrix protein 1 (ECM1), transcript variant 1, mRNA
NM 004418	Homo sapiens dual specificity phosphatase 2 (DUSP2), mRNA
NM_004961	Homo sapiens gamma-aminobutyric acid (GABA) A receptor, epsilon (GABRE), transcript variant 1, mRNA
NM_021990	Homo sapiens gamma-aminobutyric acid (GABA) A receptor, epsilon (GABRE), transcript variant 4, mRNA
NM_021987	Homo sapiens gamma-aminobutyric acid (GABA) A receptor, epsilon (GABRE), transcript variant 3, mRNA
NM_021984	Homo sapiens gamma-aminobutyric acid (GABA) A receptor, epsilon (GABRE), transcript variant 2, mRNA
NM_004090	Homo sapiens dual specificity phosphatase 3 (vaccinia virus phosphatase VH1-related) (DUSP3), mRNA
NM_001398	Homo sapiens enoyl Coenzyme A hydratase 1, peroxisomal (ECH1), mRNA
NM_001946	Homo sapiens dual specificity phosphatase 6 (DUSP6), transcript variant 1, mRNA
NM_001952	Homo sapiens E2F transcription factor 6 (E2F6), mRNA
NM 001950	Homo sapiens E2F transcription factor 4, p107/p130-binding (E2F4), mRNA
NM 001949	Homo sapiens E2F transcription factor 3 (E2F3) mRNA, complete cds
NM_005225	Homo sapiens E2F transcription factor 1 (E2F1), mRNA
NM_022977	Homo sapiens fatty-acid-Coenzyme A ligase, long-chain 4 (FACL4), transcript variant 2, mRNA
NM_004457	Homo sapiens fatty-acid-Coenzyme A ligase, long-chain 3 (FACL3), mRNA
NM 021122	Homo sapiens fatty-acid-Coenzyme A ligase, long-chain 2 (FACL2), mRNA
NM 002473	Homo sapiens myosin, heavy polypeptide 9, non-muscle (MYH9), mRNA
NM_001926	Homo sapiens defensin, alpha 6, Paneth cell-specific (DEFA6), mRNA
NM 005217	Homo sapiens defensin, alpha 3, neutrophil-specific (DEFA3), mRNA
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NM_021911 Homo sapiens gamma-aminobutyric acid (GABA) A receptor, beta 3 (GABRB3), transcript variant 2, mRNA		
NM_021611 Home sapiens gamma-aminobutyric acid (GABA) A receptor, beta 2 (GABRB2), transcript variant 1, mRNA Home sapiens gamma-aminobutyric acid (GABA) A receptor, beta 3 (GABRB3), transcript variant 1, mRNA Home sapiens gamma-aminobutyric acid (GABA) A receptor, beta 1 (GABRB1), mRNA Home sapiens gamma-aminobutyric acid (GABA) A receptor, beta 1 (GABRB1), mRNA MM_022650 Home sapiens RAS p21 protein activator (GTPase activating protein) 1 (RASA1), transcript variant 2, mRNA NM_022377 Home sapiens intercellular adhesion molecule 5, telencephalin (ICAM5), mRNA NM_022377 Home sapiens intercellular adhesion molecule 4, Landsteiner-Wiener blood group (ICAM4), transcript variant 1, mRNA NM_001544 Home sapiens intercellular adhesion molecule 4, Landsteiner-Wiener blood group (ICAM4), transcript variant 1, mRNA NM_002162 Home sapiens intercellular adhesion molecule 3 (ICAM3), mRNA NM_002308 Home sapiens intercellular adhesion molecule 2 (ICAM2), mRNA NM_022307 Home sapiens islet cell autoantigen 1 (69kD) (ICA1), transcript variant 3, mRNA NM_022307 Home sapiens islet cell autoantigen 1 (69kD) (ICA1), transcript variant 1, mRNA NM_022578 Home sapiens chorionic somatomammotropin hormone-like 1 (CSHL1), transcript variant 3, mRNA Home sapiens chorionic somatomammotropin hormone-like 1 (CSHL1), transcript variant 3, mRNA Home sapiens chorionic somatomammotropin hormone-like 1 (CSHL1), transcript variant 3, mRNA NM_022579 Home sapiens chorionic somatomammotropin hormone-like 1 (CSHL1), transcript variant 3, mRNA Home sapiens chorionic somatomammotropin hormone-like 1 (CSHL1), transcript variant 3, mRNA Home sapiens chorionic somatomammotropin hormone 2 (CSH2), transcript variant 4, mRNA Home sapiens chorionic somatomammotropin hormone 2 (CSH2), transcript variant 1, mRNA Home sapiens chorionic somatomammotropin hormone 1 (placental lactogen) (CSH1), transcript variant 1, mRNA Home sapiens chorionic somatomammotropin hormone 1 (placental lactogen) (CSH1), tran	NM_021912	
NM_000812 Homo sapiens gamma-aminobutyric acid (GABA) A receptor, beta 3 (GABR3), transcript variant 1, mRNA Homo sapiens gamma-aminobutyric acid (GABA) A receptor, beta 1 (GABRB1), mRNA Homo sapiens RAS p21 protein activator (GTPase activating protein) 1 (RASA1), transcript variant 2, mRNA MM_03259 Homo sapiens intercellular adhesion molecule 5, telencephalin (ICAM5), mRNA NM_022377 Homo sapiens intercellular adhesion molecule 4, Landsteiner-Wiener blood group (ICAM4), transcript variant 2, mRNA NM_001544 Homo sapiens intercellular adhesion molecule 4, Landsteiner-Wiener blood group (ICAM4), transcript variant 1, mRNA Homo sapiens intercellular adhesion molecule 3 (ICAM3), mRNA NM_002162 Homo sapiens intercellular adhesion molecule 2 (ICAM2), mRNA NM_002330 Homo sapiens intercellular adhesion molecule 2 (ICAM2), mRNA NM_002330 Homo sapiens islet cell autoantigen 1 (69kD) (ICA1), transcript variant 3, mRNA Homo sapiens islet cell autoantigen 1 (69kD) (ICA1), transcript variant 1, mRNA NM_022307 Homo sapiens chorionic somatomammotropin hormone-like 1 (CSHL1), transcript variant 5, mRNA Homo sapiens chorionic somatomammotropin hormone-like 1 (CSHL1), transcript variant 4, mRNA Homo sapiens chorionic somatomammotropin hormone-like 1 (CSHL1), transcript variant 1, mRNA Homo sapiens chorionic somatomammotropin hormone-like 1 (CSHL1), transcript variant 2, mRNA Homo sapiens chorionic somatomammotropin hormone-like 1 (CSHL1), transcript variant 1, mRNA Homo sapiens chorionic somatomammotropin hormone 2 (CSH2), transcript variant 3, mRNA Homo sapiens chorionic somatomammotropin hormone 2 (CSH2), transcript variant 3, mRNA Homo sapiens chorionic somatomammotropin hormone 1 (placental lactogen) (CSH1), transcript variant 3, mRNA Homo sapiens chorionic somatomammotropin hormone 1 (placental lactogen) (CSH1), transcript variant 3, mRNA Homo sapiens chorionic somatomammotropin hormone 1 (placental lactogen) (CSH1), transcript variant 3, mRNA Homo sapiens chor	NM_021911	Homo sapiens gamma-aminobutyric acid (GABA) A receptor, beta 2
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RASA1), transcript variant 2, mRNA	NM_000812	Homo sapiens gamma-aminobutyric acid (GABA) A receptor, beta 1
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NM_022377 Homo sapiens intercellular adhesion molecule 4, Landsteiner-Wiener blood group (ICAM4), transcript variant 2, mRNA	NM 003259	Homo sapiens intercellular adhesion molecule 5, telencephalin (ICAM5), mRNA
NM_002162 Homo sapiens intercellular adhesion molecule 4, Landsteiner-Wiener blood group (ICAM4), transcript variant 1, mRNA	NM_022377	Homo sapiens intercellular adhesion molecule 4, Landsteiner-Wiener blood
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NM_022439 Homo sapiens mal, T-cell differentiation protein (MAL), transcript variant c,	_	Homo sapiens mal, T-cell differentiation protein (MAL), transcript variant d,
	NM_022439	Homo sapiens mal, T-cell differentiation protein (MAL), transcript variant c,

	mRNA
NM_022438	Homo sapiens mal, T-cell differentiation protein (MAL), transcript variant b,
	mRNA
NM_001790	Homo sapiens cell division cycle 25C (CDC25C), transcript variant 1, mRNA
NM_022809	Homo sapiens cell division cycle 25C (CDC25C), transcript variant 2, mRNA
NM_021141	Homo sapiens X-ray repair complementing defective repair in Chinese hamster
	cells 5 (double-strand-break rejoining; Ku autoantigen, 80kD) (XRCC5), mRNA
NM_022550	Homo sapiens X-ray repair complementing defective repair in Chinese hamster
	cells 4 (XRCC4), transcript variant 3, mRNA
NM_022406	Homo sapiens X-ray repair complementing defective repair in Chinese hamster
	cells 4 (XRCC4), transcript variant 2, mRNA
NM_005432	Homo sapiens X-ray repair complementing defective repair in Chinese hamster
	cells 3 (XRCC3), mRNA
NM_003401	Homo sapiens X-ray repair complementing defective repair in Chinese hamster
	cells 4 (XRCC4), transcript variant 1, mRNA
NM_022405	Homo sapiens X transporter protein 3 (XT3), transcript variant 2, mRNA
NM_016192	Homo sapiens transmembrane protein with EGF-like and two follistatin-like
	domains 2 (TMEFF2), mRNA
NM_006786	Homo sapiens urotensin 2 (UTS2), transcript variant 2, mRNA
NM_021995	Homo sapiens urotensin 2 (UTS2), transcript variant 1, mRNA
NM_003353	Homo sapiens urocortin (UCN), mRNA
NM_021991	Homo sapiens junction plakoglobin (JUP), transcript variant 2, mRNA
NM_021737	Homo sapiens chloride channel 6 (CLCN6), transcript variant ClC-6d, mRNA
NM_021736	Homo sapiens chloride channel 6 (CLCN6), transcript variant ClC-6c, mRNA
NM_021735	Homo sapiens chloride channel 6 (CLCN6), transcript variant ClC-6b, mRNA
NM_006536	Homo sapiens chloride channel, calcium activated, family member 2 (CLCA2),
	mRNA
NM_004000	Homo sapiens chitinase 3-like 2 (CHI3L2), mRNA
NM_002641	Homo sapiens phosphatidylinositol glycan, class A (paroxysmal nocturnal
	hemoglobinuria) (PIGA), transcript variant 1, mRNA
NM_020473	Homo sapiens phosphatidylinositol glycan, class A (paroxysmal nocturnal
	hemoglobinuria) (PIGA), transcript variant 3, mRNA
NM_020472	Homo sapiens phosphatidylinositol glycan, class A (paroxysmal nocturnal
_	hemoglobinuria) (PIGA), transcript variant 2, mRNA
NM_001699	Homo sapiens AXL receptor tyrosine kinase (AXL), transcript variant 2, mRNA
NM_021913	Homo sapiens AXL receptor tyrosine kinase (AXL), transcript variant 1, mRNA
NM_016188	Homo sapiens actin-like 6 (ACTL6), mRNA
NM_000509	Homo sapiens fibrinogen, gamma polypeptide (FGG), transcript variant gamma-
	A, mRNA
NM_021870	Homo sapiens fibrinogen, gamma polypeptide (FGG), transcript variant gamma-
	B, mRNA
NM_005141	Homo sapiens fibrinogen, B beta polypeptide (FGB), mRNA
NM_021871	Homo sapiens fibrinogen, A alpha polypeptide (FGA), transcript variant alpha,
	mRNA
NM_000508	Homo sapiens fibrinogen, A alpha polypeptide (FGA), transcript variant alpha-E,
_	mRNA
NM 000920	Homo sapiens pyruvate carboxylase (PC), nuclear gene encoding mitochondrial
	protein, transcript variant A, mRNA
NM 022172	Homo sapiens pyruvate carboxylase (PC), nuclear gene encoding mitochondrial
	protein, transcript variant 2, mRNA
NM 004358	Homo sapiens cell division cycle 25B (CDC25B), transcript variant 1, mRNA
NM 021874	Homo sapiens cell division cycle 25B (CDC25B), transcript variant 4, mRNA
	CD Case), authority variant 7, mid in

NM_001832	Homo sapiens colipase, pancreatic (CLPS), mRNA
NM_021795	Homo sapiens ELK4, ETS-domain protein (SRF accessory protein 1) (ELK4),
	transcript variant b, mRNA
NM_021709	Homo sapiens CD27-binding (Siva) protein (SIVA), transcript variant 2, mRNA
NM_006427	Homo sapiens CD27-binding (Siva) protein (SIVA), transcript variant 1, mRNA
NM_021804	Homo sapiens angiotensin I converting enzyme (peptidyl-dipeptidase A) 2 (ACE2), mRNA
NM_020208	Homo sapiens X transporter protein 3 (XT3), transcript variant 1, mRNA
NM_021030	Homo sapiens zinc finger protein 14 (KOX 6) (ZNF14), mRNA
NM_020485	Homo sapiens Rhesus blood group, CcEe antigens (RHCE), mRNA
NM_016232	Homo sapiens interleukin 1 receptor-like 1 (IL1RL1), mRNA
NM_001680	Homo sapiens FXYD domain-containing ion transport regulator 2 (FXYD2),
L	transcript variant a, mRNA
NM_021603	Homo sapiens FXYD domain-containing ion transport regulator 2 (FXYD2), transcript variant b, mRNA
NM 005387	Homo sapiens nucleoporin 98kD (NUP98), mRNA
NM 021602	Homo sapiens CD79B antigen (immunoglobulin-associated beta) (CD79B),
	transcript variant 2, mRNA
NM_000626	Homo sapiens CD79B antigen (immunoglobulin-associated beta) (CD79B), transcript variant 1, mRNA
NM_021601	Homo sapiens CD79A antigen (immunoglobulin-associated alpha) (CD79A),
_	transcript variant 2, mRNA
NM_021599	Homo sapiens a disintegrin-like and metalloprotease (reprolysin type) with
	thrombospondin type 1 motif, 2 (ADAMTS2), transcript variant 2, mRNA
NM_006988	Homo sapiens a disintegrin-like and metalloprotease (reprolysin type) with
	thrombospondin type 1 motif, 1 (ADAMTS1), mRNA
NM_004069	Homo sapiens adaptor-related protein complex 2, sigma 1 subunit (AP2S1), transcript variant AP17, mRNA
NM 021575	Homo sapiens adaptor-related protein complex 2, sigma 1 subunit (AP2S1),
_	transcript variant AP17delta, mRNA
NM_021574	Homo sapiens breakpoint cluster region (BCR), transcript variant 2, mRNA
NM 004327	Homo sapiens breakpoint cluster region (BCR), transcript variant 1, mRNA
NM 007327	Homo sapiens glutamate receptor, ionotropic, N-methyl D-aspartate 1 (GRIN1),
_	transcript variant NR1-3, mRNA
NM_021569	Homo sapiens glutamate receptor, ionotropic, N-methyl D-aspartate 1 (GRIN1),
	transcript variant NR1-2, mRNA
NM_020984	Homo sapiens choline acetyltransferase (CHAT), transcript variant R, mRNA
NM_020985	Homo sapiens choline acetyltransferase (CHAT), transcript variant N1, mRNA
NM_020549	Homo sapiens choline acetyltransferase (CHAT), transcript variant M, mRNA
NM_001615	Homo sapiens actin, gamma 2, smooth muscle, enteric (ACTG2), mRNA
NM_020986	Homo sapiens choline acetyltransferase (CHAT), transcript variant N2, mRNA
NM_018662	Homo sapiens disrupted in schizophrenia 1 (DISC1), mRNA
NM_018406	Homo sapiens mucin 4, tracheobronchial (MUC4), mRNA
NM 017783	Homo sapiens hypothetical protein FLJ20357 (FLJ20357), mRNA
NM_004532	Homo sapiens mucin 4, tracheobronchial (MUC4), mRNA
NM_012215	Homo sapiens meningioma expressed antigen 5 (hyaluronidase) (MGEA5), mRNA
NM_020326	Homo sapiens ATP-binding cassette, sub-family D (ALD), member 4 (ABCD4),
2.111_020020	transcript variant 5, mRNA
NM_020325	Homo sapiens ATP-binding cassette, sub-family D (ALD), member 4 (ABCD4),
- 1111_020020	
NM_020324	transcript variant 4, mRNA Homo sapiens ATP-binding cassette, sub-family D (ALD), member 4 (ABCD4),

	transarint variant 2 mPNA
ND4 020222	transcript variant 3, mRNA Homo sapiens ATP-binding cassette, sub-family D (ALD), member 4 (ABCD4),
NM_020323	transcript variant 2, mRNA
NR (020209	Homo sapiens ATP-binding cassette, sub-family C (CFTR/MRP), member 9
NM_020298	
NT 000007	(ABCC9), transcript variant SUR2A-delta-14, mRNA
NM_020297	Homo sapiens ATP-binding cassette, sub-family C (CFTR/MRP), member 9
NR 001070	(ABCC9), transcript variant SUR2B, mRNA
NM_021270	Homo sapiens leukocyte-associated Ig-like receptor 2 (LAIR2), transcript variant
NR 6 000000	2, mRNA
NM_002288	Homo sapiens leukocyte-associated Ig-like receptor 2 (LAIR2), transcript variant
NM 020983	1, mRNA Homo sapiens adenylate cyclase 6 (ADCY6), transcript variant 2, mRNA
NM_015270	Homo sapiens adenylate cyclase 6 (ADCY6), transcript variant 1, mRNA
NM_020987	Homo sapiens ankyrin 3, node of Ranvier (ankyrin G) (ANK3), transcript variant 1, mRNA
NM_020977	Homo sapiens ankyrin 2, neuronal (ANK2), transcript variant 2, mRNA
NM_001148	Homo sapiens ankyrin 2, neuronal (ANK2), transcript variant 1, mRNA
NM_020481	Homo sapiens ankyrin 1, erythrocytic (ANK1), transcript variant 8, mRNA
NM_020480	Homo sapiens ankyrin 1, erythrocytic (ANK1), transcript variant 7, mRNA
NM_020479	Homo sapiens ankyrin 1, erythrocytic (ANK1), transcript variant 6, mRNA
NM_020478	Homo sapiens ankyrin 1, erythrocytic (ANK1), transcript variant 5, mRNA
NM_020477	Homo sapiens ankyrin 1, erythrocytic (ANK1), transcript variant 2, mRNA
NM_000037	Homo sapiens ankyrin 1, erythrocytic (ANK1), transcript variant 3, mRNA
NM_020476	Homo sapiens ankyrin 1, erythrocytic (ANK1), transcript variant 1, mRNA
NM_020475	Homo sapiens ankyrin 1, erythrocytic (ANK1), transcript variant 4, mRNA
NM_021056	Homo sapiens tuberous sclerosis 2 (TSC2), transcript variant 3, mRNA
NM_021055	Homo sapiens tuberous sclerosis 2 (TSC2), transcript variant 2, mRNA
NM_000548	Homo sapiens tuberous sclerosis 2 (TSC2), transcript variant 1, mRNA
NM_004041	Homo sapiens arrestin, beta 1 (ARRB1), transcript variant 1, mRNA
NM_020251	Homo sapiens arrestin, beta 1 (ARRB1), transcript variant 2, mRNA
NM_000872	Homo sapiens 5-hydroxytryptamine (serotonin) receptor 7 (adenylate cyclase-
_	coupled) (HTR7), transcript variant a, mRNA
NM_019860	Homo sapiens 5-hydroxytryptamine (serotonin) receptor 7 (adenylate cyclase-
	coupled) (HTR7), transcript variant b, mRNA
NM_019859	Homo sapiens 5-hydroxytryptamine (serotonin) receptor 7 (adenylate cyclase-
	coupled) (HTR7), transcript variant d, mRNA
NM_004228	Homo sapiens pleckstrin homology, Sec7 and coiled/coil domains 2 (cytohesin-
	2) (PSCD2), transcript variant 2, mRNA
NM_017457	Homo sapiens pleckstrin homology, Sec7 and coiled/coil domains 2 (cytohesin-
	2) (PSCD2), transcript variant 1, mRNA
NM_004302	Homo sapiens activin A receptor, type IB (ACVR1B), transcript variant 1,
	mRNA
NM_020328	Homo sapiens activin A receptor, type IB (ACVR1B), transcript variant 3,
	mRNA
NM_020327	Homo sapiens activin A receptor, type IB (ACVR1B), transcript variant 2, mRNA
NM_012082	Homo sapiens Friend of GATA2 (FOG2), mRNA
NM 000578	Homo sapiens solute carrier family 11 (proton-coupled divalent metal ion
_	transporters), member 1 (SLC11A1), mRNA
NM_021094	Homo sapiens solute carrier family 21 (organic anion transporter), member 3
_	(SLC21A3), mRNA
NM_003739	Homo sapiens aldo-keto reductase family 1, member C3 (3-alpha hydroxysteroid
	1

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37.5 000505	dehydrogenase, type II) (AKR1C3), mRNA
NM_000735	Homo sapiens glycoprotein hormones, alpha polypeptide (CGA), mRNA
NM_014272	Homo sapiens a disintegrin-like and metalloprotease (reprolysin type) with
	thrombospondin type 1 motif, 7 (ADAMTS7), mRNA
NM_019863	Homo sapiens coagulation factor VIII, procoagulant component (hemophilia A)
	(F8), transcript variant 2, mRNA
NM_000132	Homo sapiens coagulation factor VIII, procoagulant component (hemophilia A)
	(F8), transcript variant 1, mRNA
NM_019616	Homo sapiens coagulation factor VII (serum prothrombin conversion
	accelerator) (F7), transcript variant 2, mRNA
NM_000131	Homo sapiens coagulation factor VII (serum prothrombin conversion
	accelerator) (F7), transcript variant 1, mRNA
NM_007219	Homo sapiens ring finger protein 24 (RNF24), mRNA
NM_021010	Homo sapiens defensin, alpha 5, Paneth cell-specific (DEFA5), mRNA
NM_016250	Homo sapiens N-myc downstream-regulated gene 2 (NDRG2), mRNA
NM_020525	Homo sapiens interleukin 22 (IL22), mRNA
NM_006774	Homo sapiens indolethylamine N-methyltransferase (INMT), mRNA
NM_014310	Homo sapiens similar to mouse Ras, dexamethasone-induced 1 (RASD1), mRNA
NM_020322	Homo sapiens amiloride-sensitive cation channel 3, testis (ACCN3), transcript
1111_020322	variant 3, mRNA
NM_020321	Homo sapiens amiloride-sensitive cation channel 3, testis (ACCN3), transcript
	variant 2, mRNA
NM_020334	Homo sapiens a disintegrin and metalloproteinase domain 30 (ADAM30),
	transcript variant 2, mRNA
NM_019559	Homo sapiens coagulation factor XI (plasma thromboplastin antecedent) (F11),
	transcript variant 2, mRNA
NM_000128	Homo sapiens coagulation factor XI (plasma thromboplastin antecedent) (F11),
_	transcript variant 1, mRNA
NM_000443	Homo sapiens ATP-binding cassette, sub-family B (MDR/TAP), member 4
	(ABCB4), transcript variant A, mRNA
NM_018850	Homo sapiens ATP-binding cassette, sub-family B (MDR/TAP), member 4
	(ABCB4), transcript variant C, mRNA
NM_018849	Homo sapiens ATP-binding cassette, sub-family B (MDR/TAP), member 4
	(ABCB4), transcript variant B, mRNA
NM_020038	Homo sapiens ATP-binding cassette, sub-family C (CFTR/MRP), member 3
	(ABCC3), transcript variant MRP3B, mRNA
NM_020037	Homo sapiens ATP-binding cassette, sub-family C (CFTR/MRP), member 3
	(ABCC3), transcript variant MRP3A, mRNA
NM_003786	Homo sapiens ATP-binding cassette, sub-family C (CFTR/MRP), member 3
	(ABCC3), transcript variant MRP3, mRNA
NM_019624	Homo sapiens ATP-binding cassette, sub-family B (MDR/TAP), member 9
	(ABCB9), transcript variant 2, mRNA
NM_019625	Homo sapiens ATP-binding cassette, sub-family B (MDR/TAP), member 9
-	(ABCB9), transcript variant 1, mRNA
NM_004996	Homo sapiens ATP-binding cassette, sub-family C (CFTR/MRP), member 1
	(ABCC1), transcript variant 1, mRNA
NM_019902	Homo sapiens ATP-binding cassette, sub-family C (CFTR/MRP), member 1
	(ABCC1), transcript variant 7, mRNA
NM_019901	Homo sapiens ATP-binding cassette, sub-family C (CFTR/MRP), member 1
	(ABCC1), transcript variant 6, mRNA
NM 019900	Homo sapiens ATP-binding cassette, sub-family C (CFTR/MRP), member 1
	,,

	(ABCC1), transcript variant 5, mRNA
NM 019899	Homo sapiens ATP-binding cassette, sub-family C (CFTR/MRP), member 1
1111_017077	(ABCC1), transcript variant 4, mRNA
NM 019898	Homo sapiens ATP-binding cassette, sub-family C (CFTR/MRP), member 1
	(ABCC1), transcript variant 3, mRNA
NM 019862	Homo sapiens ATP-binding cassette, sub-family C (CFTR/MRP), member 1
, , , , , , , , , , , , , , , , , , , ,	(ABCC1), transcript variant 2, mRNA
NM 019903	Homo sapiens adducin 3 (gamma) (ADD3), transcript variant 2, mRNA
NM 001640	Homo sapiens N-acylaminoacyl-peptide hydrolase (APEH), mRNA
NM 019858	Homo sapiens protein A (A), transcript variant A-2, mRNA
NM 000407	Homo sapiens glycoprotein Ib (platelet), beta polypeptide (GP1BB), mRNA
NM 015675	Homo sapiens growth arrest and DNA-damage-inducible, beta (GADD45B),
- 1.1.1_0 1.2 0 1.0	mRNA
NM 016824	Homo sapiens adducin 3 (gamma) (ADD3), transcript variant 1, mRNA
NM 020039	Homo sapiens amiloride-sensitive cation channel 2, neuronal (ACCN2),
-	transcript variant 1, mRNA
NM_005388	Homo sapiens phosducin-like (PDCL), mRNA
NM_017585	Homo sapiens solute carrier family 2 (facilitated glucose transporter), member 6
	(SLC2A6), mRNA
NM_020238	Homo sapiens inner centromere protein antigens (135kD, 155kD) (INCENP),
	mRNA
NM_006908	Homo sapiens ras-related C3 botulinum toxin substrate 1 (rho family, small GTP
	binding protein Rac1) (RAC1), transcript variant Rac1, mRNA
NM_018890	Homo sapiens ras-related C3 botulinum toxin substrate 1 (rho family, small GTP
\\\\\\\\\\\\\\\\\\\\\\\\\\\\\\\\\\\\\\	binding protein Rac1) (RAC1), transcript variant Rac1b, mRNA
NM_018891	Homo sapiens laminin, gamma 2 (nicein (100kD), kalinin (105kD), BM600
	(100kD), Herlitz junctional epidermolysis bullosa)) (LAMC2), transcript variant
NM_013430	2, mRNA Homo sapiens gamma-glutamyltransferase 1 (GGT1), transcript variant 3,
14141_013430	mRNA
NM_013421	Homo sapiens gamma-glutamyltransferase 1 (GGT1), transcript variant 2,
1111_015421	mRNA
NM 004954	Homo sapiens ELKL motif kinase (EMK1), transcript variant 2, mRNA
NM_017490	Homo sapiens ELKL motif kinase (EMK1), transcript variant 1, mRNA
NM 004105	Homo sapiens EGF-containing fibulin-like extracellular matrix protein 1
	(EFEMP1), transcript variant 1, mRNA
NM_002403	Homo sapiens microfibrillar-associated protein 2 (MFAP2), transcript variant 2,
	mRNA
NM_017459	Homo sapiens microfibrillar-associated protein 2 (MFAP2), transcript variant 1,
	mRNA
NM_005115	Homo sapiens major vault protein (MVP), transcript variant 2, mRNA
NM_017458	Homo sapiens major vault protein (MVP), transcript variant 1, mRNA
NM_018894	Homo sapiens EGF-containing fibulin-like extracellular matrix protein 1
	(EFEMP1), transcript variant 2, mRNA
NM_016519	Homo sapiens ameloblastin, enamel matrix protein (AMBN), mRNA
NM_017492	Homo sapiens ataxin 2 related protein (A2LP), transcript variant 2, mRNA
NM_007193	Homo sapiens annexin A10 (ANXA10), mRNA
NM_019102	Homo sapiens homeo box A5 (HOXA5), mRNA
NM_018971	Homo sapiens G protein-coupled receptor 27 (GPR27), mRNA
NM_003379	Homo sapiens villin 2 (ezrin) (VIL2), mRNA
NM_016830	Homo sapiens vesicle-associated membrane protein 1 (synaptobrevin 1)
<u></u>	(VAMP1), transcript variant VAMP-1B, mRNA

NM_014231	Homo sapiens vesicle-associated membrane protein 1 (synaptobrevin 1)
	(VAMP1), transcript variant VAMP-1A, mRNA
NM_017489	Homo sapiens telomeric repeat binding factor (NIMA-interacting) 1 (TERF1), transcript variant 1, mRNA
NM 003218	Homo sapiens telomeric repeat binding factor (NIMA-interacting) 1 (TERF1),
14141_003218	transcript variant 2, mRNA
NM_017455	Homo sapiens stromal cell derived factor receptor 1 (SDFR1), transcript variant
	alpha, mRNA
NM_007098	Homo sapiens clathrin, heavy polypeptide-like 1 (CLTCL1), transcript variant 2,
	mRNA
NM_017451	Homo sapiens BAI1-associated protein 2 (BAIAP2), transcript variant 2, mRNA
NM_017450	Homo sapiens BAI1-associated protein 2 (BAIAP2), transcript variant 1, mRNA
NM_001617_	Homo sapiens adducin 2 (beta) (ADD2), transcript variant beta-1, mRNA
NM_017488	Homo sapiens adducin 2 (beta) (ADD2), transcript variant beta-4, mRNA
NM_017487	Homo sapiens adducin 2 (beta) (ADD2), transcript variant beta-6b, mRNA
NM_017486	Homo sapiens adducin 2 (beta) (ADD2), transcript variant beta-6a, mRNA
NM_017485	Homo sapiens adducin 2 (beta) (ADD2), transcript variant beta-5a, mRNA
NM_017484	Homo sapiens adducin 2 (beta) (ADD2), transcript variant beta-3b, mRNA
NM_017483	Homo sapiens adducin 2 (beta) (ADD2), transcript variant beta-3a, mRNA
NM_017482	Homo sapiens adducin 2 (beta) (ADD2), transcript variant beta-2, mRNA
NM 018561	Homo sapiens DKFZP586D2223 protein (DKFZP586D2223), mRNA
NM 018413	Homo sapiens chondroitin 4-sulfotransferase (C4ST), mRNA
NM 017835	Homo sapiens chromosome 21 open reading frame 59 (C21ORF59), mRNA
NM 018226	Homo sapiens arginyl aminopeptidase (aminopeptidase B)-like 1 (RNPEPL1),
	mRNA
NM_018204	Homo sapiens cytoskeleton associated protein 2 (CKAP2), mRNA
NM_018200	Homo sapiens high-mobility group 20A (HMG20A), mRNA
NM_017595	Homo sapiens I-kappa-B-interacting Ras-like protein 2 (KBRAS2), mRNA
NM_017613	Homo sapiens downstream neighbor of SON (DONSON), mRNA
NM_017596	Homo sapiens KIAA0449 protein (KIAA0449), mRNA
NM_017456	Homo sapiens pleckstrin homology, Sec7 and coiled/coil domains 1(cytohesin 1)
	(PSCD1), transcript variant 2, mRNA
NM_016829	Homo sapiens 8-oxoguanine DNA glycosylase (OGG1), nuclear gene encoding
	mitochondrial protein, transcript variant 2e, mRNA
NM_016828	Homo sapiens 8-oxoguanine DNA glycosylase (OGG1), nuclear gene encoding
	mitochondrial protein, transcript variant 2d, mRNA
NM_016827	Homo sapiens 8-oxoguanine DNA glycosylase (OGG1), nuclear gene encoding
	mitochondrial protein, transcript variant 2c, mRNA
NM_016826	Homo sapiens 8-oxoguanine DNA glycosylase (OGG1), nuclear gene encoding
	mitochondrial protein, transcript variant 2b, mRNA
NM_016821	Homo sapiens 8-oxoguanine DNA glycosylase (OGG1), nuclear gene encoding
	mitochondrial protein, transcript variant 2a, mRNA
NM_016820	Homo sapiens 8-oxoguanine DNA glycosylase (OGG1), nuclear gene encoding
	mitochondrial protein, transcript variant 1c, mRNA
NM_016819	Homo sapiens 8-oxoguanine DNA glycosylase (OGG1), nuclear gene encoding
	mitochondrial protein, transcript variant 1b, mRNA
NM_002197	Homo sapiens aconitase 1, soluble (ACO1), mRNA
NM_016841	Homo sapiens microtubule-associated protein tau (MAPT), transcript variant 4,
	mRNA
NM_016835	Homo sapiens microtubule-associated protein tau (MAPT), transcript variant 1,
	mRNA
NM_016834	Homo sapiens microtubule-associated protein tau (MAPT), transcript variant 3,

	mRNA
NM 016938	Homo sapiens EGF-containing fibulin-like extracellular matrix protein 2
	(EFEMP2), mRNA
NM 005569	Homo sapiens LIM domain kinase 2 (LIMK2), transcript variant 2a, mRNA
NM 016733	Homo sapiens LIM domain kinase 2 (LIMK2), transcript variant 2b, mRNA
NM_002314	Homo sapiens LIM domain kinase 1 (LIMK1), transcript variant 1, mRNA
NM 016735	Homo sapiens LIM domain kinase 1 (LIMK1), transcript variant dLIMK, mRNA
NM 006855	Homo sapiens KDEL (Lys-Asp-Glu-Leu) endoplasmic reticulum protein
	retention receptor 3 (KDELR3), transcript variant 1, mRNA
NM_016657	Homo sapiens KDEL (Lys-Asp-Glu-Leu) endoplasmic reticulum protein
_	retention receptor 3 (KDELR3), transcript variant 2, mRNA
NM_002101	Homo sapiens glycophorin C (Gerbich blood group) (GYPC), transcript variant
_	1, mRNA
NM_016815	Homo sapiens glycophorin C (Gerbich blood group) (GYPC), transcript variant
	2, mRNA
NM_005242	Homo sapiens coagulation factor II (thrombin) receptor-like 1 (F2RL1), mRNA
NM_016818	Homo sapiens ATP-binding cassette, sub-family G (WHITE), member 1
	(ABCG1), transcript variant 2, mRNA
NM_004915	Homo sapiens ATP-binding cassette, sub-family G (WHITE), member 1
ND 6 000515	(ABCG1), transcript variant 1, mRNA
NM_002542	Homo sapiens 8-oxoguanine DNA glycosylase (OGG1), nuclear gene encoding
NT 6 000665	mitochondrial protein, transcript variant 1a, mRNA
NM_000665	Homo sapiens acetylcholinesterase (YT blood group) (ACHE), transcript variant
NIM 012000	E4-E6, mRNA Home serious mesenchume homes her 1 (MEOV1) transaciut varient 2 mRNA
NM_013999 NM_003927	Homo sapiens mesenchyme homeo box 1 (MEOX1), transcript variant 2, mRNA Homo sapiens methyl-CpG binding domain protein 2 (MBD2), transcript variant
7.41AT_00237	1, mRNA
NM_015832	Homo sapiens methyl-CpG binding domain protein 2 (MBD2), transcript variant
	testis-specific, mRNA
NM_002384	Homo sapiens methyl-CpG binding domain protein 1 (MBD1), transcript variant
	4, mRNA
NM_015847	Homo sapiens methyl-CpG binding domain protein 1 (MBD1), transcript variant
	PCM1, mRNA
NM_015846	Homo sapiens methyl-CpG binding domain protein 1 (MBD1), transcript variant
·	1, mRNA
NM_015845	Homo sapiens methyl-CpG binding domain protein 1 (MBD1), transcript variant
	2, mRNA
NM_015844	Homo sapiens methyl-CpG binding domain protein 1 (MBD1), transcript variant
NB (000011	3, mRNA
NM_002311	Homo sapiens ligase III, DNA, ATP-dependent (LIG3), transcript variant beta, mRNA
NIM 012075	
NM_013975	Homo sapiens ligase III, DNA, ATP-dependent (LIG3), transcript variant alpha, mRNA
NM 014190	Homo sapiens adducin 1 (alpha) (ADD1), transcript variant 3, mRNA
NM 014189	Homo sapiens adducin 1 (alpha) (ADD1), transcript variant 3, mRNA Homo sapiens adducin 1 (alpha) (ADD1), transcript variant 2, mRNA
NM_001119	Homo sapiens adducin 1 (alpha) (ADD1), transcript variant 2, mRNA Homo sapiens adducin 1 (alpha) (ADD1), transcript variant 1, mRNA
NM 015831	Homo sapiens acetylcholinesterase (YT blood group) (ACHE), transcript variant
144-015051	E4-E5, mRNA
NM 016572	Homo sapiens ubiquitin specific protease 21 (USP21), mRNA
NM_016388	Homo sapiens T-cell receptor interacting molecule (TRIM), mRNA
NM_016272	Homo sapiens transducer of ERBB2, 2 (TOB2), mRNA
NM 016135	Homo sapiens transcription factor ets (TEL2), mRNA
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NM_016247	Homo sapiens interphotoreceptor matrix proteoglycan 200 (SPACRCAN),
	mRNA
NM_016334	Homo sapiens putative G-protein coupled receptor (SH120), mRNA
NM_016124	Homo sapiens Rhesus blood group, D antigen (RHD), mRNA
NM_015865	Homo sapiens solute carrier family 14 (urea transporter), member 1 (Kidd blood group) (SLC14A1), mRNA
NM_016112	Homo sapiens polycystic kidney disease 2-like 1 (PKD2L1), mRNA
NM_016318	Homo sapiens purinergic receptor P2X, ligand-gated ion channel, 2 (P2RX2), mRNA
NM_016653	Homo sapiens sterile-alpha motif and leucine zipper containing kinase AZK (ZAK), mRNA
NM_016556	Homo sapiens GT198, complete ORF (HUMGT198A), mRNA
NM_016431	Homo sapiens mitogen-activated protein kinase 8 interacting protein 2 (MAPK8IP2), mRNA
NM_016377	Homo sapiens A kinase (PRKA) anchor protein 7 (AKAP7), mRNA
NM_016346	Homo sapiens nuclear receptor subfamily 2, group E, member 3 (NR2E3), mRNA
NM_016325	Homo sapiens zinc finger protein 274 (ZNF274), mRNA
NM_016324	Homo sapiens zinc finger protein 274 (ZNF274), mRNA
NM_016293	Homo sapiens bridging integrator 2 (BIN2), mRNA
NM_015909	Homo sapiens neuroblastoma-amplified protein (LOC51594), mRNA
NM_015890	Homo sapiens spondyloepiphyseal dysplasia, late, pseudogene (SEDLP), mRNA
NM_015885	Homo sapiens PCF11p homolog (PCF11), mRNA
NM_015991	Homo sapiens complement component 1, q subcomponent, alpha polypeptide
ND (016201	(C1QA), mRNA
NM_016201 NM_016157	Homo sapiens Leman coiled-coil protein (LCCP), mRNA
NM_015869	Homo sapiens trophinin (TRO), mRNA
	Homo sapiens peroxisome proliferative activated receptor, gamma (PPARG), mRNA
NM_016615	Homo sapiens solute carrier family 6 (neurotransmitter transporter, GABA), member 13 (SLC6A13), mRNA
NM 016389	Homo sapiens NS1-binding protein (NS1-BP), mRNA
NM 016648	Homo sapiens HDCMA18P protein (HDCMA18P), mRNA
NM_016527	Homo sapiens hydroxyacid oxidase 2 (long chain) (HAO2), mRNA
NM_016263	Homo sapiens Fzr1 protein (FZR1), mRNA
NM_016602	Homo sapiens G protein-coupled receptor 2 (GPR2), mRNA
NM_015892	Homo sapiens B cell RAG associated protein (BRAG), mRNA
NM_016187	Homo sapiens bridging integrator 2 (BIN2), mRNA
NM_003373	Homo sapiens vinculin (VCL), transcript variant VCL, mRNA
NM_014000	Homo sapiens vinculin (VCL), transcript variant meta-VCL, mRNA
NM_013992	Homo sapiens paired box gene 8 (PAX8), transcript variant PAX8E, mRNA
NM_013988	Homo sapiens Parkinson disease (autosomal recessive, juvenile) 2, parkin
ND4 012005	(PARK2), transcript variant 3, mRNA
NM_013987	Homo sapiens Parkinson disease (autosomal recessive, juvenile) 2, parkin (PARK2), transcript variant 2, mRNA
NM_013985	
NM 013984	Homo sapiens neuregulin 2 (NRG2), transcript variant 6, mRNA
NM 013983	Homo sapiens neuregulin 2 (NRG2), transcript variant 5, mRNA
NM 013982	Homo sapiens neuregulin 2 (NRG2), transcript variant 4, mRNA
NM_013981	Homo sapiens neuregulin 2 (NRG2), transcript variant 3, mRNA
NM_013964	Homo sapiens neuregulin 2 (NRG2), transcript variant 2, mRNA
NM_013962	Homo sapiens neuregulin 1 (NRG1), transcript variant HRG-alpha, mRNA
1111 013302	Homo sapiens neuregulin 1 (NRG1), transcript variant GGF2, mRNA

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NM_013961	Homo sapiens neuregulin 1 (NRG1), transcript variant GGF, mRNA
NM_013960	Homo sapiens neuregulin 1 (NRG1), transcript variant ndf43, mRNA
NM_013959	Homo sapiens neuregulin 1 (NRG1), transcript variant SMDF, mRNA
NM_013958	Homo sapiens neuregulin 1 (NRG1), transcript variant HRG-beta3, mRNA
NM_013957	Homo sapiens neuregulin 1 (NRG1), transcript variant HRG-beta2, mRNA
NM_013956	Homo sapiens neuregulin 1 (NRG1), transcript variant HRG-beta1, mRNA
NM_013955	Homo sapiens NADPH oxidase 1 (NOX1), transcript variant NOH-1Lv, mRNA
NM_013954	Homo sapiens NADPH oxidase 1 (NOX1), transcript variant NOH-1S, mRNA
NM_013995	Homo sapiens lysosomal-associated membrane protein 2 (LAMP2), transcript variant LAMP2B, mRNA
NM_007334	Homo sapiens killer cell lectin-like receptor subfamily D, member 1 (KLRD1), transcript variant 2, mRNA
NM_002262	Homo sapiens killer cell lectin-like receptor subfamily D, member 1 (KLRD1), transcript variant 1, mRNA
NM_013976	Homo sapiens glutaryl-Coenzyme A dehydrogenase (GCDH), nuclear gene encoding mitochondrial protein, transcript variant 2, mRNA
NM_015841	Homo sapiens adenosine deaminase, RNA-specific (ADAR), transcript variant
1111_015071	ADAR-c, mRNA
NM_015840	Homo sapiens adenosine deaminase, RNA-specific (ADAR), transcript variant ADAR-b, mRNA
NM 001111	Homo sapiens adenosine deaminase, RNA-specific (ADAR), transcript variant
_	ADAR-a, mRNA
NM_014925	Homo sapiens KIAA1002 protein (KIAA1002), mRNA
NM_014905	Homo sapiens glutaminase (GLS), mRNA
NM_014833	Homo sapiens KIAA0618 gene product (KIAA0618), mRNA
NM_014863	Homo sapiens B cell RAG associated protein (BRAG), mRNA
NM_015646	Homo sapiens RAP1B, member of RAS oncogene family (RAP1B), mRNA
NM_015423	Homo sapiens aminoadipate-semialdehyde dehydrogenase-phosphopantetheinyl transferase (AASDHPPT), mRNA
NM_015523	Homo sapiens small fragment nuclease (DKFZP566E144), mRNA
NM_014397	Homo sapiens NIMA (never in mitosis gene a)-related kinase 6 (NEK6), mRNA
NM_014249	Homo sapiens nuclear receptor subfamily 2, group E, member 3 (NR2E3), mRNA
NM 014361	Homo sapiens contactin 5 (CNTN5), mRNA
NM_014341	Homo sapiens mitochondrial carrier homolog 1 (MTCH1), nuclear gene
	encoding mitochondrial protein, mRNA
NM_014556	Homo sapiens Ellis van Creveld syndrome (EVC), mRNA
NM_014306	Homo sapiens hypothetical protein (HSPC117), mRNA
NM_014593	Homo sapiens CpG binding protein (CGBP), mRNA
NM_014567	Homo sapiens breast cancer anti-estrogen resistance 1 (BCAR1), mRNA
NM_014273	Homo sapiens a disintegrin-like and metalloprotease (reprolysin type) with thrombospondin type 1 motif, 6 (ADAMTS6), mRNA
NM_014244	Homo sapiens a disintegrin-like and metalloprotease (reprolysin type) with thrombospondin type 1 motif, 2 (ADAMTS2), transcript variant 1, mRNA
NM_014449	Homo sapiens protein A (A), transcript variant A-1, mRNA
NM_007319	Homo sapiens presenilin 1 (Alzheimer disease 3) (PSEN1), transcript variant I-374., mRNA
NM_007318	Homo sapiens presenilin 1 (Alzheimer disease 3) (PSEN1), transcript variant I-463, mRNA
NM_013953	Homo sapiens paired box gene 8 (PAX8), transcript variant PAX8D, mRNA
NM_013952	Homo sapiens paired box gene 8 (PAX8), transcript variant PAX8C, mRNA
NM 013951	
	Homo sapiens paired box gene 8 (PAX8), transcript variant PAX8B, mRNA

NM_013945	Homo sapiens paired box gene 7 (PAX7), transcript variant 2, mRNA
NM_013942	Homo sapiens paired box gene 3 (Waardenburg syndrome 1) (PAX3), transcript
	variant PAX3B, mRNA
NM_013411	Homo sapiens adenylate kinase 2 (AK2), nuclear gene encoding mitochondrial
	protein, transcript variant AK2B, mRNA
NM_000631	Homo sapiens neutrophil cytosolic factor 4 (40kD) (NCF4), transcript variant 1,
	mRNA
NM_013416	Homo sapiens neutrophil cytosolic factor 4 (40kD) (NCF4), transcript variant 2,
	mRNA
NM_006125	Homo sapiens Rho GTPase activating protein 6 (ARHGAP6), transcript variant
\	3, mRNA
NM_013427	Homo sapiens Rho GTPase activating protein 6 (ARHGAP6), transcript variant
ND 6 012402	1, mRNA
NM_013423	Homo sapiens Rho GTPase activating protein 6 (ARHGAP6), transcript variant
ND4 012422	4, mRNA
NM_013422	Homo sapiens Rho GTPase activating protein 6 (ARHGAP6), transcript variant
NM_001174	5, mRNA
11111_001174	Homo sapiens Rho GTPase activating protein 6 (ARHGAP6), transcript variant 2, mRNA
NM 013436	Homo sapiens NCK-associated protein 1 (NCKAP1), mRNA
NM 012310	Homo sapiens kinesin family member 4A (KIF4A), mRNA
NM 013449	Homo sapiens bromodomain adjacent to zinc finger domain, 2A (BAZ2A),
11212_015 1 15	mRNA
NM_007333	Homo sapiens killer cell lectin-like receptor subfamily C, member 3 (KLRC3),
-	transcript variant NKG2-H, mRNA
NM 007328	Homo sapiens killer cell lectin-like receptor subfamily C, member 1 (KLRC1),
	transcript variant NKG2-B, mRNA
NM_002259	Homo sapiens killer cell lectin-like receptor subfamily C, member 1 (KLRC1),
	transcript variant NKG2-A, mRNA
NM_004214	Homo sapiens fibroblast growth factor (acidic) intracellular binding protein
	(FIBP), mRNA
NM_006350	Homo sapiens follistatin (FST), transcript variant FST317, mRNA
NM_013409	Homo sapiens follistatin (FST), transcript variant FST344, mRNA
NM_013324	Homo sapiens cytokine inducible SH2-containing protein (CISH), mRNA
NM_012486	Homo sapiens presenilin 2 (Alzheimer disease 4) (PSEN2), transcript variant 2,
ND4 012495	mRNA
NM_012485	Homo sapiens hyaluronan-mediated motility receptor (RHAMM) (HMMR),
NM 012484	transcript variant 2, mRNA
14141_012404	Homo sapiens hyaluronan-mediated motility receptor (RHAMM) (HMMR), transcript variant 1, mRNA
NM 012483	
NM 006433	Homo sapiens granulysin (GNLY), transcript variant 519, mRNA
NM 001930	Homo sapiens granulysin (GNLY), transcript variant NKG5, mRNA
NM 013407	Homo sapiens deoxylypusine synthase (DHPS), transcript variant 1, mRNA
NM_013406	Homo sapiens deoxylypusine synthase (DHPS), transcript variant 3, mRNA
NM 013229	Homo sapiens deoxyhypusine synthase (DHPS), transcript variant 2, mRNA Homo sapiens apoptotic protease activating factor (APAF1), transcript variant 1,
	mRNA
NM 013251	Homo sapiens tachykinin 3 (neuromedin K, neurokinin beta) (TAC3), mRNA
NM 013396	Homo sapiens ubiquitin specific protease 25 (USP25), mRNA
NM 013255	Homo sapiens muskelin 1, intracellular mediator containing kelch motifs
	(MKLN1), mRNA
NM_013290	Homo sapiens GT198, complete ORF (HUMGT198A), mRNA
	7, 1111 1111 1111 1111 1111

NM_005102	Homo sapiens fasciculation and elongation protein zeta 2 (zygin II) (FEZ2), mRNA
NM_004830	Homo sapiens cofactor required for Sp1 transcriptional activation, subunit 3 (130kD) (CRSP3), mRNA
NM_009588	Homo sapiens lymphotoxin beta (TNF superfamily, member 3) (LTB), transcript variant 2, mRNA
NM_013227	Homo sapiens aggrecan 1 (chondroitin sulfate proteoglycan 1, large aggregating proteoglycan, antigen identified by monoclonal antibody A0122) (AGC1), transcript variant 2, mRNA
NM_012475	Homo sapiens ubiquitin specific protease 21 (USP21), mRNA
NM_012428	Homo sapiens stromal cell derived factor receptor 1 (SDFR1), transcript variant beta, mRNA
NM_012226	Homo sapiens purinergic receptor P2X, ligand-gated ion channel, 2 (P2RX2), mRNA
NM_012369	Homo sapiens olfactory receptor, family 2, subfamily F, member 1 (OR2F1), mRNA
NM_012218	Homo sapiens interleukin enhancer binding factor 3, 90kD (ILF3), mRNA
NM_012324	Homo sapiens mitogen-activated protein kinase 8 interacting protein 2 (MAPK8IP2), mRNA
NM_012405	Homo sapiens isoprenylcysteine carboxyl methyltransferase (ICMT), mRNA
NM_012070	Homo sapiens attractin (ATRN), mRNA
NM_006874	Homo sapiens E74-like factor 2 (ets domain transcription factor) (ELF2), mRNA
NM_007308	Homo sapiens synuclein, alpha (non A4 component of amyloid precursor) (SNCA), transcript variant NACP112, mRNA
NM_000345	Homo sapiens synuclein, alpha (non A4 component of amyloid precursor) (SNCA), transcript variant NACP140, mRNA
NM 009589	Homo sapiens arylsulfatase D (ARSD), transcript variant 2, mRNA
NM_001158	Homo sapiens amine oxidase, copper containing 2 (retina-specific) (AOC2), transcript variant 1, mRNA
NM_005910	Homo sapiens microtubule-associated protein tau (MAPT), transcript variant 2, mRNA
NM_007338	Homo sapiens deleted in lung and esophageal cancer 1 (DLEC1), transcript variant DLEC1-L1, mRNA
NM_007337	Homo sapiens deleted in lung and esophageal cancer 1 (DLEC1), transcript variant DLEC1-S3, mRNA
NM_007336	Homo sapiens deleted in lung and esophageal cancer 1 (DLEC1), transcript variant DLEC1-S2, mRNA
NM_007335	Homo sapiens deleted in lung and esophageal cancer 1 (DLEC1), transcript variant DLEC1-S1, mRNA
NM_005106	Homo sapiens deleted in lung and esophageal cancer 1 (DLEC1), transcript variant DLEC1-N1, mRNA
NM_005002	Homo sapiens NADH dehydrogenase (ubiquinone) 1 alpha subcomplex, 9 (39kD) (NDUFA9), mRNA
NM_003771	Homo sapiens keratin, hair, acidic, 6 (KRTHA6), mRNA
NM_000438	Homo sapiens paired box gene 3 (Waardenburg syndrome 1) (PAX3), transcript variant PAX3A, mRNA
NM_007052	Homo sapiens NADPH oxidase 1 (NOX1), transcript variant NOH-1L, mRNA
NM_006715	Homo sapiens mannosidase, alpha, class 2C, member 1 (MAN2C1), mRNA
NM_007325	Homo sapiens glutamate receptor, ionotrophic, AMPA 3 (GRIA3), transcript variant flip, mRNA
NM_005813	Homo sapiens protein kinase C, nu (PRKCN), mRNA
NM_000398	Homo sapiens diaphorase (NADH) (cytochrome b-5 reductase) (DIA1), nuclear
	, and the second

	gene encoding mitochondrial protein, transcript variant M, mRNA
NM 007306	Homo sapiens breast cancer 1, early onset (BRCA1), transcript variant BRCA1-
MM_007300	
NR 6 000000	exon4, mRNA
NM_007305	Homo sapiens breast cancer 1, early onset (BRCA1), transcript variant BRCA1-
17.6.005004	delta9-10-11b, mRNA
NM_007304	Homo sapiens breast cancer 1, early onset (BRCA1), transcript variant BRCA1-
	deltallb, mRNA
NM_007303	Homo sapiens breast cancer 1, early onset (BRCA1), transcript variant BRCA1-
	delta11, mRNA
NM_007302	Homo sapiens breast cancer 1, early onset (BRCA1), transcript variant BRCA1-
	delta9-10, mRNA
NM_007301	Homo sapiens breast cancer 1, early onset (BRCA1), transcript variant BRCA1-
	delta15-17, mRNA
NM 007300	Homo sapiens breast cancer 1, early onset (BRCA1), transcript variant BRCA1-
_	delta14-18, mRNA
NM_007299	Homo sapiens breast cancer 1, early onset (BRCA1), transcript variant BRCA1-
_	delta14-17, mRNA
NM_007298	Homo sapiens breast cancer 1, early onset (BRCA1), transcript variant BRCA1-
	delta9-11, mRNA
NM_007297	Homo sapiens breast cancer 1, early onset (BRCA1), transcript variant BRCA1-
	delta2-10, mRNA
NM_007296	Homo sapiens breast cancer 1, early onset (BRCA1), transcript variant BRCA1a',
11111_007230	mRNA
NM_007295	Homo sapiens breast cancer 1, early onset (BRCA1), transcript variant BRCA1b,
1111_007255	mRNA
NM_007294	Homo sapiens breast cancer 1, early onset (BRCA1), transcript variant BRCA1a,
11111_007254	mRNA
NM_007322	Homo sapiens RAN binding protein 3 (RANBP3), transcript variant RANBP3-d,
	mRNA
NM_007321	Homo sapiens RAN binding protein 3 (RANBP3), transcript variant RANBP3-c,
11112_007521	mRNA
NM 007320	Homo sapiens RAN binding protein 3 (RANBP3), transcript variant RANBP3-b,
11112_007520	mRNA
NM_000754	Homo sapiens catechol-O-methyltransferase (COMT), transcript variant MB-
11111_000754	COMT, mRNA
NM 007310	Homo sapiens catechol-O-methyltransferase (COMT), transcript variant S-
14147_00/210	COMT, mRNA
NM 000714	
NIVI_000714	Homo sapiens benzodiazapine receptor (peripheral) (BZRP), nuclear gene
NB4 007211	encoding mitochondrial protein, transcript variant PBR, mRNA
NM_007311	Homo sapiens benzodiazapine receptor (peripheral) (BZRP), nuclear gene
ND (007214	encoding mitochondrial protein, transcript variant PBR-S, mRNA
NM_007314	Homo sapiens v-abl Abelson murine leukemia viral oncogene homolog 2 (arg,
37.6 005010	Abelson-related gene) (ABL2), transcript variant b, mRNA
NM_007313	Homo sapiens v-abl Abelson murine leukemia viral oncogene homolog 1
12000000	(ABL1), transcript variant b, mRNA
NM_005157	Homo sapiens v-abl Abelson murine leukemia viral oncogene homolog 1
L	(ABL1), transcript variant a, mRNA
NM_006325	Homo sapiens RAN, member RAS oncogene family (RAN), mRNA
NM_000902	Homo sapiens membrane metallo-endopeptidase (neutral endopeptidase,
	enkephalinase, CALLA, CD10) (MME), transcript variant 1, mRNA
NM_007289	Homo sapiens membrane metallo-endopeptidase (neutral endopeptidase,
L	enkephalinase, CALLA, CD10) (MME), transcript variant 2b, mRNA

NM_007288	Homo sapiens membrane metallo-endopeptidase (neutral endopeptidase, enkephalinase, CALLA, CD10) (MME), transcript variant 2a, mRNA
NM_007287	Homo sapiens membrane metallo-endopeptidase (neutral endopeptidase,
27.6.006404	enkephalinase, CALLA, CD10) (MME), transcript variant 1bis, mRNA
NM_006481	Homo sapiens transcription factor 2, hepatic; LF-B3; variant hepatic nuclear factor (TCF2), transcript variant b, mRNA
NM_006884	Homo sapiens short stature homeobox 2 (SHOX2), transcript variant SHOX2a, mRNA
NM_003030	Homo sapiens short stature homeobox 2 (SHOX2), transcript variant SHOX2b, mRNA
NM_003005	Homo sapiens selectin P (granule membrane protein 140kD, antigen CD62) (SELP), mRNA
NM_006718	Homo sapiens pleiomorphic adenoma gene-like 1 (PLAGL1), transcript variant 2, mRNA
NM_005888	Homo sapiens solute carrier family 25 (mitochondrial carrier; phosphate carrier),
	member 3 (SLC25A3), nuclear gene encoding mitochondrial protein, transcript variant 1a, mRNA
NM_006491	Homo sapiens neuro-oncological ventral antigen 1 (NOVA1), transcript variant
	3, mRNA
NM_006489	Homo sapiens neuro-oncological ventral antigen 1 (NOVA1), transcript variant
	2, mRNA
NM_007088	Homo sapiens calbindin 2, (29kD, calretinin) (CALB2), transcript variant
_	CALB2c, mRNA
NM 007087	Homo sapiens calbindin 2, (29kD, calretinin) (CALB2), transcript variant
_	CALB2b, mRNA
NM 001740	Homo sapiens calbindin 2, (29kD, calretinin) (CALB2), transcript variant
_	CALB2, mRNA
NM_007292	Homo sapiens acyl-Coenzyme A oxidase 1, palmitoyl (ACOX1), transcript variant 2, mRNA
NM_004035	Homo sapiens acyl-Coenzyme A oxidase 1, palmitoyl (ACOX1), transcript variant 1, mRNA
NM_000632	Homo sapiens integrin, alpha M (complement component receptor 3, alpha; also
	known as CD11b (p170), macrophage antigen alpha polypeptide) (ITGAM),
NIM 007007	mRNA Home senions elethric light nelsonatide (Leh) (CLTP) mPNA
NM_007097 NM_007099	Homo sapiens clathrin, light polypeptide (Lcb) (CLTB), mRNA
	Homo sapiens acid phosphatase 1, soluble (ACP1), transcript variant b, mRNA
NM_007177	Homo sapiens TU3A protein (TU3A), mRNA
NM 007245	Homo sapiens ataxin 2 related protein (A2LP), transcript variant 1, mRNA
NM_006487	Homo sapiens fibulin 1 (FBLN1), transcript variant A, mRNA
NM_006486	Homo sapiens fibulin 1 (FBLN1), transcript variant D, mRNA
NM_006485	Homo sapiens fibulin 1 (FBLN1), transcript variant B, mRNA
NM_006721	Homo sapiens adenosine kinase (ADK), transcript variant ADK-long, mRNA
NM_006132	Homo sapiens bone morphogenetic protein 1 (BMP1), transcript variant BMP1-4, mRNA
NM_006131	Homo sapiens bone morphogenetic protein 1 (BMP1), transcript variant BMP1-5, mRNA
NM_006130	Homo sapiens bone morphogenetic protein 1 (BMP1), transcript variant BMP1-6, mRNA
NM_006129	Homo sapiens bone morphogenetic protein 1 (BMP1), transcript variant BMP1-3, mRNA
NM_006128	Homo sapiens bone morphogenetic protein 1 (BMP1), transcript variant BMP1-2, mRNA
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NM 002516	Homo sapiens neuro-oncological ventral antigen 2 (NOVA2), mRNA
	/
NM_007008	Homo sapiens reticulon 4 (RTN4), mRNA
NM_007046	Homo sapiens elastin microfibril interface located protein (EMILIN), mRNA
NM_007037	Homo sapiens a disintegrin-like and metalloprotease (reprolysin type) with
)D (007020	thrombospondin type 1 motif, 8 (ADAMTS8), mRNA
NM_007038	Homo sapiens a disintegrin-like and metalloprotease (reprolysin type) with
NR 6 00 6500	thrombospondin type 1 motif, 5 (aggrecanase-2) (ADAMTS5), mRNA
NM_006799	Homo sapiens protease, serine, 21 (testisin) (PRSS21), mRNA
NM_006814	Homo sapiens proteasome (prosome, macropain) inhibitor subunit 1 (PI31) (PSMF1), mRNA
NM_003466	Homo sapiens paired box gene 8 (PAX8), transcript variant PAX8A, mRNA
NM_006790	Homo sapiens titin immunoglobulin domain protein (myotilin) (TTID), mRNA
NM_006782	Homo sapiens zinc finger protein-like 1 (ZFPL1), mRNA
NM_006795	Homo sapiens EH domain containing 1 (EHD1), mRNA
NM_006588	Homo sapiens sulfotransferase family, cytosolic, 1C, member 2 (SULT1C2), mRNA
NM_006694	Homo sapiens jumping translocation breakpoint (JTB), mRNA
NM_006597	Homo sapiens heat shock 70kD protein 8 (HSPA8), mRNA
NM 006708	Homo sapiens glyoxalase I (GLO1), mRNA
NM 006703	Homo sapiens nudix (nucleoside diphosphate linked moiety X)-type motif 3
-	(NUDT3), mRNA
NM 000655	Homo sapiens selectin L (lymphocyte adhesion molecule 1) (SELL), mRNA
NM 006488	Homo sapiens ketohexokinase (fructokinase) (KHK), transcript variant b, mRNA
NM_006297	Homo sapiens X-ray repair complementing defective repair in Chinese hamster
	cells 1 (XRCC1), mRNA
NM 006339	Homo sapiens high-mobility group 20B (HMG20B), mRNA
NM 006469	Homo sapiens NS1-binding protein (NS1-BP), mRNA
NM 006340	Homo sapiens BAI1-associated protein 2 (BAIAP2), transcript variant 3, mRNA
NM_001353	Homo sapiens aldo-keto reductase family 1, member C1 (dihydrodiol dehydrogenase 1; 20-alpha (3-alpha)-hydroxysteroid dehydrogenase) (AKR1C1), mRNA
NM_000202	Homo sapiens iduronate 2-sulfatase (Hunter syndrome) (IDS), transcript variant 1, mRNA
NM_005890	Homo sapiens growth arrest-specific 7 (GAS7), transcript variant b, mRNA
NM_006123	Homo sapiens iduronate 2-sulfatase (Hunter syndrome) (IDS), transcript variant 2, mRNA
NM_006053	Homo sapiens T-cell, immune regulator 1 (TCIRG1), mRNA
NM_005990	Homo sapiens serine/threonine kinase 10 (STK10), mRNA
NM_006019	Homo sapiens T-cell, immune regulator 1 (TCIRG1), mRNA
NM_006041	Homo sapiens heparan sulfate (glucosamine) 3-O-sulfotransferase 3B1 (HS3ST3B1), mRNA
NM_006042	Homo sapiens heparan sulfate (glucosamine) 3-O-sulfotransferase 3A1 (HS3ST3A1), mRNA
NM_006043	Homo sapiens heparan sulfate (glucosamine) 3-O-sulfotransferase 2 (HS3ST2), mRNA
NM_000557	Homo sapiens growth differentiation factor 5 (cartilage-derived morphogenetic protein-1) (GDF5), mRNA
NM_005847	Homo sapiens solute carrier family 23 (nucleobase transporters), member 2 (SLC23A2), mRNA
NM_005751	Homo sapiens A kinase (PRKA) anchor protein (yotiao) 9 (AKAP9), mRNA
NM_005691	Homo sapiens ATP-binding cassette, sub-family C (CFTR/MRP), member 9 (ABCC9), transcript variant SUR2A, mRNA

NM_005688	Homo sapiens ATP-binding cassette, sub-family C (CFTR/MRP), member 5 (ABCC5), mRNA
NM_005730	Homo sapiens conserved gene amplified in osteosarcoma (OS4), mRNA
NM_005562	Homo sapiens laminin, gamma 2 (nicein (100kD), kalinin (105kD), BM600
	(100kD), Herlitz junctional epidermolysis bullosa)) (LAMC2), transcript variant 1, mRNA
NM_005534	Homo sapiens interferon gamma receptor 2 (interferon gamma transducer 1) (IFNGR2), mRNA
NM_005682	Homo sapiens G protein-coupled receptor 56 (GPR56), mRNA
NM_005666	Homo sapiens H factor (complement)-like 3 (HFL3), mRNA
NM_005503	Homo sapiens amyloid beta (A4) precursor protein-binding, family A, member 2 (X11-like) (APBA2), mRNA
NM_005431	Homo sapiens X-ray repair complementing defective repair in Chinese hamster cells 2 (XRCC2), mRNA
NM_005465	Homo sapiens v-akt murine thymoma viral oncogene homolog 3 (protein kinase B, gamma) (AKT3), mRNA
NM_005446	Homo sapiens purinergic receptor P2X-like 1, orphan receptor (P2RXL1), mRNA
NM_005336	Homo sapiens high density lipoprotein binding protein (vigilin) (HDLBP), mRNA
NM_005265	Homo sapiens gamma-glutamyltransferase 1 (GGT1), transcript variant 1, mRNA
NM_005243	Homo sapiens Ewing sarcoma breakpoint region 1 (EWSR1), transcript variant EWS, mRNA
NM_005236	Homo sapiens excision repair cross-complementing rodent repair deficiency, complementation group 4 (ERCC4), mRNA
NM_005075	Homo sapiens solute carrier family 21 (organic anion transporter), member 3 (SLC21A3), mRNA
NM_005050	Homo sapiens ATP-binding cassette, sub-family D (ALD), member 4 (ABCD4), transcript variant 1, mRNA
NM_005006	Homo sapiens NADH dehydrogenase (ubiquinone) Fe-S protein 1 (75kD) (NADH-coenzyme Q reductase) (NDUFS1), mRNA
NM_005135	Homo sapiens solute carrier family 12 (potassium/chloride transporters), member 6 (SLC12A6), mRNA
NM_004968	Homo sapiens islet cell autoantigen 1 (69kD) (ICA1), transcript variant 2, mRNA
NM_005114	Homo sapiens heparan sulfate (glucosamine) 3-O-sulfotransferase 1 (HS3ST1), mRNA
NM_004958	Homo sapiens FK506 binding protein 12-rapamycin associated protein 1 (FRAP1), mRNA
NM_001478	Homo sapiens UDP-N-acetyl-alpha-D-galactosamine:(N-acetylneuraminyl)-galactosylglucosylceramide N-acetylgalactosaminyltransferase (GalNAc-T) (GALGT), mRNA
NM_004031	Homo sapiens interferon regulatory factor 7 (IRF7), transcript variant d, mRNA
NM_004030	Homo sapiens interferon regulatory factor 7 (IRF7), transcript variant c, mRNA
NM_004029	Homo sapiens interferon regulatory factor 7 (IRF7), transcript variant b, mRNA
NM_004034	Homo sapiens annexin A7 (ANXA7), transcript variant 2, mRNA
NM_001156	Homo sapiens annexin A7 (ANXA7), transcript variant 1, mRNA
NM_004033	Homo sapiens annexin A6 (ANXA6), transcript variant 2, mRNA
NM_001155	Homo sapiens annexin A6 (ANXA6), transcript variant 1, mRNA
NM 004629	Homo sapiens Fanconi anemia, complementation group G (FANCG), mRNA
NM_004738	Homo sapiens VAMP (vesicle-associated membrane protein)-associated protein

Homo sapiens paired box gene 6 (aniridia, keratitis) (PAX6), mRNA
Homo sapiens natriuretic peptide receptor B/guanylate cyclase B (atrionatriuretic
peptide receptor B) (NPR2), mRNA
Homo sapiens KIT ligand (KITLG), mRNA
Homo sapiens transferrin (TF), mRNA
Homo sapiens paired box gene 2 (PAX2), transcript variant e, mRNA
Homo sapiens paired box gene 2 (PAX2), transcript variant d, mRNA
Homo sapiens paired box gene 2 (PAX2), transcript variant c, mRNA
Homo sapiens paired box gene 2 (PAX2), transcript variant a, mRNA
Homo sapiens paired box gene 2 (PAX2), transcript variant b, mRNA
Homo sapiens ketohexokinase (fructokinase) (KHK), transcript variant a, mRNA
Homo sapiens endothelin receptor type B (EDNRB), transcript variant 1, mRNA
Homo sapiens carnitine acetyltransferase (CRAT), nuclear gene encoding
mitochondrial protein, transcript variant mitochondrial, mRNA
Homo sapiens CDC-like kinase 3 (CLK3), transcript variant phclk3/152, mRNA
Homo sapiens CDC-like kinase 2 (CLK2), transcript variant phclk2/139, mRNA
Homo sapiens adaptor-related protein complex 2, beta 1 subunit (AP2B1), mRNA
Homo sapiens chromodomain helicase DNA binding protein 3 (CHD3), mRNA
Homo sapiens chromosome condensation 1-like (CHC1L), mRNA
Homo sapiens CD3Z antigen, zeta polypeptide (TiT3 complex) (CD3Z), mRNA
Homo sapiens B-cell CLL/lymphoma 2 (BCL2), nuclear gene encoding
mitochondrial protein, transcript variant beta, mRNA
Homo sapiens B-cell CLL/lymphoma 2 (BCL2), nuclear gene encoding
mitochondrial protein, transcript variant alpha, mRNA
Homo sapiens butyrylcholinesterase (BCHE), mRNA
Homo sapiens transcription termination factor, RNA polymerase II (TTF2), mRNA
Homo sapiens tumor protein 63 kDa with strong homology to p53 (TP63), mRNA
Homo sapiens interleukin 1 receptor-like 1 (IL1RL1), mRNA
Homo sapiens sex determining region Y (SRY), mRNA
Homo sapiens solute carrier family 4, sodium bicarbonate cotransporter, member 7 (SLC4A7), mRNA
Homo sapiens solute carrier family 4, sodium bicarbonate cotransporter, member 4 (SLC4A4), mRNA
Homo sapiens secretin receptor (SCTR), mRNA
Homo sapiens RAS p21 protein activator (GTPase activating protein) 1 (RASA1), transcript variant 1, mRNA
Homo sapiens RAN binding protein 3 (RANBP3), transcript variant RANBP3-a, mRNA
Homo sapiens proteasome (prosome, macropain) 26S subunit, non-ATPase, 13 (PSMD13), mRNA
Homo sapiens presenilin 2 (Alzheimer disease 4) (PSEN2), transcript variant 1, mRNA
Homo sapiens presenilin 1 (Alzheimer disease 3) (PSEN1), transcript variant I-467, mRNA
Homo sapiens procollagen (type III) N-endopeptidase (PCOLN3), mRNA
Homo sapiens mitogen-activated protein kinase 9 (MAPK9), mRNA
Homo sapiens pleiomorphic adenoma gene-like 1 (PLAGL1), transcript variant 1, mRNA
Homo sapiens solute carrier family 25 (mitochondrial carrier; phosphate carrier),

	1 2 (01 025 42)
_	member 3 (SLC25A3), nuclear gene encoding mitochondrial protein, transcript
NR 6 000504	variant 1b, mRNA
NM_002584	Homo sapiens paired box gene 7 (PAX7), transcript variant 1, mRNA
NM_000280	Homo sapiens paired box gene 6 (aniridia, keratitis) (PAX6), mRNA
NM_002555	Homo sapiens solute carrier family 22 (organic cation transporter), member 1-like (SLC22A1L), mRNA
NM_000907	Homo sapiens natriuretic peptide receptor B/guanylate cyclase B (atrionatriuretic peptide receptor B) (NPR2), mRNA
NM_002515	Homo sapiens neuro-oncological ventral antigen 1 (NOVA1), transcript variant 1, mRNA
NM 003204	Homo sapiens nuclear factor (erythroid-derived 2)-like 1 (NFE2L1), mRNA
NM 003970	Homo sapiens myomesin (M-protein) 2 (165kD) (MYOM2), mRNA
NM 000899	Homo sapiens KIT ligand (KITLG), mRNA
NM_002394	Homo sapiens solute carrier family 3 (activators of dibasic and neutral amino acid transport), member 2 (SLC3A2), mRNA
NM_001879	Homo sapiens mannan-binding lectin serine protease 1 (C4/C2 activating component of Ra-reactive factor) (MASP1), mRNA
NM_002353	Homo sapiens tumor-associated calcium signal transducer 2 (TACSTD2), mRNA
NM_002341	Homo sapiens lymphotoxin beta (TNF superfamily, member 3) (LTB), transcript variant 1, mRNA
NM_002294	Homo sapiens lysosomal-associated membrane protein 2 (LAMP2), transcript variant LAMP2A, mRNA
NM 002264	Homo sapiens karyopherin alpha 1 (importin alpha 5) (KPNA1), mRNA
NM_002261	Homo sapiens killer cell lectin-like receptor subfamily C, member 3 (KLRC3), transcript variant NKG2-E, mRNA
NM 002230	Homo sapiens junction plakoglobin (JUP), transcript variant 1, mRNA
NM_001566	Homo sapiens inositol polyphosphate-4-phosphatase, type I, 107kD (INPP4A), transcript variant b, mRNA
NM 002164	Homo sapiens indoleamine-pyrrole 2,3 dioxygenase (INDO), mRNA
NM_003822	Homo sapiens nuclear receptor subfamily 5, group A, member 2 (NR5A2), mRNA
NM_000836	Homo sapiens glutamate receptor, ionotropic, N-methyl D-aspartate 2D (GRIN2D), mRNA
NM_000828	Homo sapiens glutamate receptor, ionotrophic, AMPA 3 (GRIA3), transcript variant flop, mRNA
NM_002056	Homo sapiens glutamine-fructose-6-phosphate transaminase 1 (GFPT1), mRNA
NM_000161	Homo sapiens GTP cyclohydrolase 1 (dopa-responsive dystonia) (GCH1), mRNA
NM_000159	Homo sapiens glutaryl-Coenzyme A dehydrogenase (GCDH), nuclear gene encoding mitochondrial protein, transcript variant 1, mRNA
NM_003644	Homo sapiens growth arrest-specific 7 (GAS7), transcript variant a, mRNA
NM_000817	Homo sapiens glutamate decarboxylase 1 (brain, 67kD) (GAD1), transcript variant GAD67, mRNA
NM_000813	Homo sapiens gamma-aminobutyric acid (GABA) A receptor, beta 2 (GABRB2), transcript variant 2, mRNA
NM 000146	Homo sapiens ferritin, light polypeptide (FTL), mRNA
NM_001996	Homo sapiens fibulin 1 (FBLN1), transcript variant C, mRNA
NM_001995	Homo sapiens fatty-acid-Coenzyme A ligase, long-chain 1 (FACL1), nuclear gene encoding mitochondrial protein, mRNA
NM_001973	Homo sapiens ELK4, ETS-domain protein (SRF accessory protein 1) (ELK4), transcript variant a, mRNA

NM_003991	Homo sapiens endothelin receptor type B (EDNRB), transcript variant 2, mRNA
NM_001925	Homo sapiens defensin, alpha 4, corticostatin (DEFA4), mRNA
NM_001359	Homo sapiens 2,4-dienoyl CoA reductase 1, mitochondrial (DECR1), nuclear
	gene encoding mitochondrial protein, mRNA
NM_001337	Homo sapiens chemokine (C-X3-C) receptor 1 (CX3CR1), mRNA
NM_001835	Homo sapiens clathrin, heavy polypeptide-like 1 (CLTCL1), transcript variant 1,
L	mRNA
NM_001834	Homo sapiens clathrin, light polypeptide (Lcb) (CLTB), transcript variant
	nonbrain, mRNA
NM_003992	Homo sapiens CDC-like kinase 3 (CLK3), transcript variant phclk3, mRNA
NM_003993	Homo sapiens CDC-like kinase 2 (CLK2), transcript variant phclk2, mRNA
NM_001286	Homo sapiens chloride channel 6 (CLCN6), transcript variant ClC-6a, mRNA
NM_001285	Homo sapiens chloride channel, calcium activated, family member 1 (CLCA1),
	mRNA
NM_001825	Homo sapiens creatine kinase, mitochondrial 2 (sarcomeric) (CKMT2), nuclear
	gene encoding mitochondrial protein, mRNA
NM_003465	Homo sapiens chitinase 1 (chitotriosidase) (CHIT1), mRNA
NM_001783	Homo sapiens CD79A antigen (immunoglobulin-associated alpha) (CD79A),
ND 6 001100	transcript variant 1, mRNA
NM_001199	Homo sapiens bone morphogenetic protein 1 (BMP1), transcript variant BMP1-
ND (001660	1, mRNA
NM_001669	Homo sapiens arylsulfatase D (ARSD), transcript variant 1, mRNA
NM_001170	Homo sapiens aquaporin 7 (AQP7), mRNA
NM_001160	Homo sapiens apoptotic protease activating factor (APAF1), transcript variant 2, mRNA
NM_001149	Homo sapiens ankyrin 3, node of Ranvier (ankyrin G) (ANK3), transcript variant
	2, mRNA
NM_001625	Homo sapiens adenylate kinase 2 (AK2), nuclear gene encoding mitochondrial
	protein, transcript variant AK2A, mRNA
NM_001135	Homo sapiens aggrecan 1 (chondroitin sulfate proteoglycan 1, large aggregating
	proteoglycan, antigen identified by monoclonal antibody A0122) (AGC1),
ND 6 001100	transcript variant 1, mRNA
NM_001123	Homo sapiens adenosine kinase (ADK), transcript variant ADK-short, mRNA
NM_003812	Homo sapiens a disintegrin and metalloproteinase domain 23 (ADAM23), mRNA
NM_001095	Homo sapiens amiloride-sensitive cation channel 2, neuronal (ACCN2),
	transcript variant 2, mRNA
NM_016184	Homo sapiens C-type (calcium dependent, carbohydrate-recognition domain)
ND4 000:00	lectin, superfamily member 6 (CLECSF6), mRNA
NM_003186	Homo sapiens transgelin (TAGLN), mRNA
NM_004084	Homo sapiens defensin, alpha 1, myeloid-related sequence (DEFA1), mRNA
NM 022908	Homo sapiens hypothetical protein FLJ12442 (FLJ12442), mRNA
NM_022906	Homo sapiens hypothetical protein FLJ13195 similar to stromal antigen 3
NIM 022002	(FLJ13195), mRNA
NM_022903	Homo sapiens hypothetical protein FLJ12800 (FLJ12800), mRNA
NM_022902	Homo sapiens hypothetical protein FLJ12496 (FLJ12496), mRNA
NM_022900 NM_022895	Homo sapiens hypothetical protein FLJ21213 (FLJ21213), mRNA
NM 006997	Homo sapiens hypothetical protein FLJ12448 (FLJ12448), mRNA
	Homo sapiens transforming, acidic coiled-coil containing protein 2 (TACC2), mRNA
NM_020979	Homo sapiens adaptor protein with pleckstrin homology and src homology 2
	domains (APS), mRNA

NM_018557	Homo sapiens low density lipoprotein-related protein 1B (deleted in tumors) (LRP1B), mRNA
NM 014921	Homo sapiens lectomedin-2 (KIAA0821), mRNA
NM 014112	Homo sapiens trichorhinophalangeal syndrome I gene (TRPS1), mRNA
NM 000539	Homo sapiens rhodopsin (opsin 2, rod pigment) (retinitis pigmentosa 4,
14141_000333	autosomal dominant) (RHO), mRNA
NM 012452	Homo sapiens transmembrane activator and CAML interactor (TACI), mRNA
NM 003564	Homo sapiens transgelin 2 (TAGLN2), mRNA
NM 003632	Homo sapiens contactin associated protein 1 (CNTNAP1), mRNA
NM 006506	Homo sapiens RAS p21 protein activator 2 (RASA2), mRNA
NM_014427	Homo sapiens copine VII (CPNE7), mRNA
NM_006032	Homo sapiens copine VI (neuronal) (CPNE6), mRNA
NM_005338	Homo sapiens huntingtin interacting protein 1 (HIP1), mRNA
NM_021973	Homo sapiens heart and neural crest derivatives expressed 2 (HAND2), mRNA
NM_005339	Homo sapiens huntingtin interacting protein 2 (HIP2), mRNA
NM_021920	Homo sapiens secretin (SCT), mRNA
NM_016491	Homo sapiens mitochondrial ribosomal protein L37 (MRPL37), mRNA
NM_014211	Homo sapiens gamma-aminobutyric acid (GABA) A receptor, pi (GABRP), mRNA
NM 004658	Homo sapiens RAS protein activator like 1 (GAP1 like) (RASAL1), mRNA
NM 004807	Homo sapiens heparan sulfate 6-O-sulfotransferase (HS6ST), mRNA
NM 002622	Homo sapiens prefoldin 1 (PFDN1), mRNA
NM 005186	Homo sapiens calpain 1, (mu/I) large subunit (CAPN1), mRNA
NM 001748	Homo sapiens calpain 2, (m/II) large subunit (CAPN2), mRNA
NM 014299	Homo sapiens bromodomain-containing 4 (BRD4), mRNA
NM 007208	Homo sapiens mitochondrial ribosomal protein L3 (MRPL3), mRNA
NM 022838	Homo sapiens hypothetical protein FLJ12969 (FLJ12969), mRNA
NM 022837	Homo sapiens hypothetical protein FLJ22833 (FLJ22833), mRNA
NM 022830	Homo sapiens hypothetical protein FLJ22347 (FLJ22347), mRNA
NM_022819	Homo sapiens phospholipase A2, group IIF (PLA2G2F), mRNA
NM_020245	Homo sapiens tubby super-family protein (TUSP), mRNA
NM_020061	Homo sapiens opsin 1 (cone pigments), long-wave-sensitive (color blindness, protan) (OPN1LW), mRNA
NM_000513	Homo sapiens opsin 1 (cone pigments), medium-wave-sensitive (color blindness, deutan) (OPN1MW), mRNA
NM_001708	Homo sapiens opsin 1 (cone pigments), short-wave-sensitive (color blindness,
1411_001700	tritan) (OPN1SW), mRNA
NM_016363	Homo sapiens glycoprotein VI (platelet) (GP6), mRNA
NM_022139	Homo sapiens GDNF family receptor alpha 4 (GFRA4), mRNA
NM 002485	Homo sapiens Nijmegen breakage syndrome 1 (nibrin) (NBS1), mRNA
NM_006052	Homo sapiens Down syndrome critical region gene 3 (DSCR3), mRNA
NM_005867	Homo sapiens Down syndrome critical region gene 4 (DSCR4), mRNA
NM_005087	Homo sapiens fragile X mental retardation, autosomal homolog 1 (FXR1), mRNA
NM 004403	Homo sapiens deafness, autosomal dominant 5 (DFNA5), mRNA
NM 000433	Homo sapiens neutrophil cytosolic factor 2 (65kD, chronic granulomatous
_	disease, autosomal 2) (NCF2), mRNA
NM 000111	Homo sapiens solute carrier family 26, member 3 (SLC26A3), mRNA
NM_000044	Homo sapiens androgen receptor (dihydrotestosterone receptor; testicular
	feminization; spinal and bulbar muscular atrophy; Kennedy disease) (AR), mRNA
NM 000333	Homo sapiens spinocerebellar ataxia 7 (olivopontocerebellar atrophy with retinal

domination) (CCA7) mDNA
degeneration) (SCA7), mRNA Homo sapiens nuclear localization signal deleted in velocardiofacial syndrome
·
(NLVCF), mRNA
Homo sapiens Wiskott-Aldrich syndrome-like (WASL), mRNA
Homo sapiens N-terminal kinase-like (NTKL), mRNA
Homo sapiens interleukin 17E (IL17E), mRNA
Homo sapiens NMN adenylyltransferase; nicotinamide mononucleotide adenylyl transferase (NMNAT), mRNA
Homo sapiens likely ortholog of yeast ARV1 (ARV1), mRNA
Homo sapiens hypothetical protein FLJ23588 (FLJ23588), mRNA
Homo sapiens hypothetical protein FLJ22127 (FLJ22127), mRNA
Homo sapiens hypothetical protein FLJ12681 (FLJ12681), mRNA
Homo sapiens hypothetical protein FLJ21935 (FLJ21935), mRNA
Homo sapiens hypothetical protein FLJ23499 (FLJ23499), mRNA
Homo sapiens hypothetical protein FLJ11730 (FLJ11730), mRNA
Homo sapiens E3 ubiquitin ligase SMURF2 (SMURF2), mRNA
Homo sapiens Fanconi anemia, complementation group F (FANCF), mRNA
Homo sapiens tRNA isopentenylpyrophosphate transferase (IPT), mRNA
Homo sapiens 3'-phosphoadenosine 5'-phosphosulfate synthase 1 (PAPSS1), mRNA
Homo sapiens 3'-phosphoadenosine 5'-phosphosulfate synthase 2 (PAPSS2), mRNA
Homo sapiens procollagen-lysine, 2-oxoglutarate 5-dioxygenase 3 (PLOD3), mRNA
Homo sapiens DiGeorge syndrome critical region gene 8 (DGCR8), mRNA
Homo sapiens Wolf-Hirschhorn syndrome candidate 1 (WHSC1), mRNA
Homo sapiens Usher syndrome 2A (autosomal recessive, mild) (USH2A), mRNA
Homo sapiens Werner syndrome (WRN), mRNA
Homo sapiens Probe hTg737 (polycystic kidney disease, autosomal recessive, in) (TG737), mRNA
Homo sapiens Down syndrome critical region gene 6 (DSCR6), mRNA
Homo sapiens McKusick-Kaufman syndrome (MKKS), mRNA
Homo sapiens cat eye syndrome chromosome region, candidate 1 (CECR1), mRNA
Homo sapiens TPA inducible gene-1 (TIG-1), mRNA
Homo sapiens Down syndrome critical region gene 5 (DSCR5), mRNA
Homo sapiens Down syndrome critical region gene 1 (DSCR1), mRNA
Homo sapiens Down syndrome critical region gene 1-like 2 (DSCR1L2), mRNA
Homo sapiens sperm associated antigen 8 (SPAG8), mRNA
Homo sapiens Pseudoautosomal GTP-binding protein-like (PGPL), mRNA
Homo sapiens protease, serine, 23 (SPUVE), mRNA
Homo sapiens elastin (supravalvular aortic stenosis, Williams-Beuren syndrome)
(ELN), mRNA
Homo sapiens protease, serine, 22 (P11), mRNA
Homo sapiens phosphorylase, glycogen; muscle (McArdle syndrome, glycogen
storage disease type V) (PYGM), mRNA
Homo sapiens myelodysplasia syndrome 1 (MDS1), mRNA
Homo sapiens Sjogren syndrome antigen A2 (60kD, ribonucleoprotein
autoantigen SS-A/Ro) (SSA2), mRNA
Homo sapiens CREB binding protein (Rubinstein-Taybi syndrome) (CREBBP), mRNA

NM_000551	Homo sapiens von Hippel-Lindau syndrome (VHL), mRNA
NM_000462	Homo sapiens ubiquitin protein ligase E3A (human papilloma virus E6-
_	associated protein, Angelman syndrome) (UBE3A), mRNA
NM_001064	Homo sapiens transketolase (Wernicke-Korsakoff syndrome) (TKT), mRNA
NM_000356	Homo sapiens Treacher Collins-Franceschetti syndrome 1 (TCOF1), mRNA
NM_000455	Homo sapiens serine/threonine kinase 11 (Peutz-Jeghers syndrome) (STK11),
	mRNA
NM_002351	Homo sapiens SH2 domain protein 1A, Duncan's disease (lymphoproliferative
	syndrome) (SH2D1A), mRNA
NM_000336	Homo sapiens sodium channel, nonvoltage-gated 1, beta (Liddle syndrome) (SCNN1B), mRNA
NM 000335	Homo sapiens sodium channel, voltage-gated, type V, alpha polypeptide (long
	(electrocardiographic) QT syndrome 3) (SCN5A), mRNA
NM 000318	Homo sapiens peroxisomal membrane protein 3 (35kD, Zellweger syndrome)
	(PXMP3), mRNA
NM_000311	Homo sapiens prion protein (p27-30) (Creutzfeld-Jakob disease, Gerstmann-
_	Strausler-Scheinker syndrome, fatal familial insomnia) (PRNP), mRNA
NM 000299	Homo sapiens plakophilin 1 (ectodermal dysplasia/skin fragility syndrome)
	(PKP1), mRNA
NM_000283	Homo sapiens phosphodiesterase 6B, cGMP-specific, rod, beta (congenital
	stationary night blindness 3, autosomal dominant) (PDE6B), mRNA
NM_003731	Homo sapiens Sjogren's syndrome nuclear autoantigen 1 (SSNA1), mRNA
NM_000260	Homo sapiens myosin VIIA (Usher syndrome 1B (autosomal recessive, severe))
	(MYO7A), mRNA
NM_003720	Homo sapiens Down syndrome critical region gene 2 (DSCR2), mRNA
NM_000195	Homo sapiens Hermansky-Pudlak syndrome (HPS), mRNA
NM_000194	Homo sapiens hypoxanthine phosphoribosyltransferase 1 (Lesch-Nyhan syndrome) (HPRT1), mRNA
NM_000171	Homo sapiens glycine receptor, alpha 1 (startle disease/hyperekplexia, stiff man
_	syndrome) (GLRA1), mRNA
NM_003494	Homo sapiens dysferlin, limb girdle muscular dystrophy 2B (autosomal
	recessive) (DYSF), mRNA
NM_000081	Homo sapiens Chediak-Higashi syndrome 1 (CHS1), mRNA
NM_000052	Homo sapiens ATPase, Cu++ transporting, alpha polypeptide (Menkes
	syndrome) (ATP7A), mRNA
NM_001635	Homo sapiens amphiphysin (Stiff-Mann syndrome with breast cancer 128kD
	autoantigen) (AMPH), mRNA
NM_022663	Homo sapiens CTAGE-1 protein (CTAGE-1), mRNA
NM_022662	Homo sapiens meiotic checkpoint regulator (MCPR), mRNA
NM_022658	Homo sapiens homeo box C8 (HOXC8), mRNA
NM_000569	Homo sapiens Fc fragment of IgG, low affinity IIIa, receptor for (CD16) (FCGR3A), mRNA
NM 000802	Homo sapiens folate receptor 1 (adult) (FOLR1), transcript variant 2, mRNA
NM 006991	Homo sapiens zinc finger protein 197 (ZNF197), mRNA
NM 018946	Homo sapiens N-acetylneuraminic acid phosphate synthase; sialic acid synthase
	(SAS), mRNA
NM_003979	Homo sapiens retinoic acid induced 3 (RAI3), mRNA
NM_021785	Homo sapiens retinoic acid induced 2 (RAI2), mRNA
NM_001436	Homo sapiens fibrillarin (FBL), mRNA
NM_012151	Homo sapiens coagulation factor VIII-associated (intronic transcript) (F8A), mRNA
NM 007170	Homo sapiens testis-specific kinase 2 (TESK2), mRNA
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NM_006285	Homo sapiens testis-specific kinase 1 (TESK1), mRNA
NM_016424	Homo sapiens cisplatin resistance-associated overexpressed protein (LUC7A), mRNA
NM 012152	Homo sapiens endothelial differentiation, lysophosphatidic acid G-protein-
	coupled receptor, 7 (EDG7), mRNA
NM 007360	Homo sapiens DNA segment on chromosome 12 (unique) 2489 expressed
_	sequence (D12S2489E), mRNA
NM 004924	Homo sapiens actinin, alpha 4 (ACTN4), mRNA
NM 001102	Homo sapiens actinin, alpha 1 (ACTN1), mRNA
NM 012128	Homo sapiens chloride channel, calcium activated, family member 4 (CLCA4),
-	mRNA
NM_014551	Homo sapiens hypothetical protein 384D8_6 (384D8-2), mRNA
NM_018977	Homo sapiens neuroligin 3 (NLGN3), mRNA
NM 001103	Homo sapiens actinin, alpha 2 (ACTN2), mRNA
NM_022569	Homo sapiens N-deacetylase/N-sulfotransferase 4 (NDST4), mRNA
NM_005892	Homo sapiens formin-like (FMNL), mRNA
NM 016370	Homo sapiens RAB9-like protein (RAB9L), mRNA
NM_012135	Homo sapiens DNA segment on chromosome 6(unique) 2654 expressed
	sequence (D6S2654E), mRNA
NM_007161	Homo sapiens DNA segment on chromosome 6 (unique) 49 expressed sequence,
	NK cell triggering receptor, p30 (D6S49E), mRNA
NM_006114	Homo sapiens DNA segment on chromosome 19 (unique) 1177 expressed
	sequence (D19S1177E), mRNA
NM_006014	Homo sapiens DNA segment on chromosome X (unique) 9879 expressed
	sequence (DXS9879E), mRNA
NM_004699	Homo sapiens DNA segment on chromosome X (unique) 9928 expressed
	sequence (DXS9928E), mRNA
NM_003683	Homo sapiens DNA segment on chromosome 21 (unique) 2056 expressed
	sequence (D21S2056E), mRNA
NM_015484	Homo sapiens GCIP-interacting protein p29 (P29), mRNA
NM_013263	Homo sapiens bromodomain-containing 7 (BRD7), mRNA
NM_022157	Homo sapiens Rag C protein (GTR2), mRNA
NM_014604	Homo sapiens Tax interaction protein 1 (TIP-1), mRNA
NM_001915	Homo sapiens cytochrome b-561 (CYB561), mRNA
NM_012188	Homo sapiens forkhead box I1 (FOXI1), mRNA
NM_016148	Homo sapiens somatostatin receptor-interacting protein (SSTRIP), mRNA
NM_022482	Homo sapiens hypothetical protein FLJ21794 (FLJ21794), mRNA
NM_022493	Homo sapiens hypothetical protein FLJ21988 (FLJ21988), mRNA
NM_022489	Homo sapiens hypothetical protein FLJ22056 (FLJ22056), mRNA
NM_022485	Homo sapiens hypothetical protein FLJ22405 (FLJ22405), mRNA
NM_022464	Homo sapiens endoplasmic reticulum chaperone SIL1, homolog of yeast (SIL1), mRNA
NM_022456	Homo sapiens hypothetical protein FLJ22548 similar to gene trap PAT 12
14141_022450	(FLJ22548), mRNA
NM 022450	Homo sapiens hypothetical protein FLJ22357 similar to epidermal growth factor
11112_022750	receptor-related protein (FLJ22357), mRNA
NM_022443	Homo sapiens myeloid leukemia factor 1 (MLF1), mRNA
NM 022136	Homo sapiens SAM domain, SH3 domain and nuclear localisation signals, 1
	(SAMSN1), mRNA
NM_012217	Homo sapiens mast cell tryptase (TPSD1), mRNA
NM 020366	Homo sapiens retinitis pigmentosa GTPase regulator interacting protein 1
	(RPGRIP1), mRNA
	1

NM_016541	Homo sapiens guanine nucleotide binding protein 13, gamma (GNG13), mRNA
NM_004204	Homo sapiens phosphatidylinositol glycan, class Q (PIGQ), mRNA
NM_014946	Homo sapiens spastic paraplegia 4 (autosomal dominant; spastin) (SPG4), mRNA
NM_022146	Homo sapiens neuropeptide FF 1; RFamide-related peptide receptor (OT7T022), mRNA
NM_004885	Homo sapiens neuropeptide G protein-coupled receptor; neuropeptide FF 2 (NPGPR), mRNA
NM 002958	Homo sapiens RYK receptor-like tyrosine kinase (RYK), mRNA
NM 002931	Homo sapiens ring finger protein 1 (RING1), mRNA
NM_021111	Homo sapiens reversion-inducing-cysteine-rich protein with kazal motifs (RECK), mRNA
NM 001655	Homo sapiens archain 1 (ARCN1), mRNA
NM 016639	Homo sapiens type I transmembrane protein Fn14 (FN14), mRNA
NM 006686	Homo sapiens actin-like 7B (ACTL7B), mRNA
NM 006687	Homo sapiens actin-like 7A (ACTL7A), mRNA
NM_005856	Homo sapiens receptor (calcitonin) activity modifying protein 3 (RAMP3), mRNA
NM_005854	Homo sapiens receptor (calcitonin) activity modifying protein 2 (RAMP2), mRNA
NM_005855	Homo sapiens receptor (calcitonin) activity modifying protein 1 (RAMP1), mRNA
NM_000475	Homo sapiens nuclear receptor subfamily 0, group B, member 1 (NR0B1), mRNA
NM_005493	Homo sapiens RAN binding protein 9 (RANBP9), mRNA
NM_004634	Homo sapiens bromodomain and PHD finger containing, 1 (BRPF1), mRNA
NM_000140	Homo sapiens ferrochelatase (protoporphyria) (FECH), nuclear gene encoding mitochondrial protein, mRNA
NM_000031	Homo sapiens aminolevulinate, delta-, dehydratase (ALAD), mRNA
NM_000027	Homo sapiens aspartylglucosaminidase (AGA), mRNA
NM_000026	Homo sapiens adenylosuccinate lyase (ADSL), mRNA
NM_000025	Homo sapiens adrenergic, beta-3-, receptor (ADRB3), mRNA
NM_000020	Homo sapiens activin A receptor type II-like 1 (ACVRL1), mRNA
NM_000019	Homo sapiens acetyl-Coenzyme A acetyltransferase 1 (acetoacetyl Coenzyme A thiolase) (ACAT1), nuclear gene encoding mitochondrial protein, mRNA
NM_000018	Homo sapiens acyl-Coenzyme A dehydrogenase, very long chain (ACADVL), nuclear gene encoding mitochondrial protein, mRNA
NM_000017	Homo sapiens acyl-Coenzyme A dehydrogenase, C-2 to C-3 short chain (ACADS), nuclear gene encoding mitochondrial protein, mRNA
NM_000016	Homo sapiens acyl-Coenzyme A dehydrogenase, C-4 to C-12 straight chain (ACADM), nuclear gene encoding mitochondrial protein, mRNA
NM 000476	Homo sapiens adenylate kinase 1 (AK1), mRNA
NM 001830	Homo sapiens chloride channel 4 (CLCN4), mRNA
NM 022365	Homo sapiens hypothetical protein similar to mouse Dnajl1 (DNAJL1), mRNA
NM_022350	Homo sapiens aminopeptidase (LOC64167), mRNA
NM 022335	Homo sapiens hypothetical protein PRO2849 (PRO2849), mRNA
NM_005259	Homo sapiens growth differentiation factor 8 (GDF8), mRNA
NM_001789	Homo sapiens cell division cycle 25A (CDC25A), mRNA
NM_022006	Homo sapiens FXYD domain-containing ion transport regulator 7 (FXYD7), mRNA
NM_022003	Homo sapiens FXYD domain-containing ion transport regulator 6 (FXYD6), mRNA

NM_020655	Homo sapiens junctophilin 3 (JPH3), mRNA
NM_002855	Homo sapiens poliovirus receptor-related 1 (herpesvirus entry mediator C;
	nectin) (PVRL1), mRNA
NM 012340	Homo sapiens nuclear factor of activated T-cells, cytoplasmic, calcineurin-
_	dependent 2 (NFATC2), mRNA
NM 006599	Homo sapiens nuclear factor of activated T-cells 5, tonicity-resonsive (NFAT5),
_	mRNA
NM 006162	Homo sapiens nuclear factor of activated T-cells, cytoplasmic, calcineurin-
	dependent 1 (NFATC1), mRNA
NM 022061	Homo sapiens ribosomal protein L17 isolog (LOC63875), mRNA
NM_022095	Homo sapiens hypothetical C2H2 zinc finger protein FLJ22504 (FLJ22504),
	mRNA
NM 022091	Homo sapiens dJ467N11.1 protein (DJ467N11.1), mRNA
NM 022084	Homo sapiens hypothetical protein dJ102H19.4 (DJ102H19.4), mRNA
NM 022077	Homo sapiens hypothetical protein dJ1141E15.2 (DJ1141E15.2), mRNA
NM 022098	Homo sapiens hypothetical protein LOC63929 (LOC63929), mRNA
NM 022081	Homo sapiens hypothetical protein bK1048E9.5 (BK1048E9.5), mRNA
NM 021081	Homo sapiens growth hormone releasing hormone (GHRH), mRNA
	Homo sapiens melanoma differentiation associated protein-5 (MDA5), mRNA
NM_022168	↓
NM_022165	Homo sapiens Lin-7b protein (LIN-7B), mRNA
NM_022161	Homo sapiens livin inhibitor-of-apotosis (LIVIN), mRNA
NM_022159	Homo sapiens ETL protein (ETL), mRNA
NM_022156	Homo sapiens PP3111 protein (PP3111), mRNA
NM_022151	Homo sapiens MAP-1 protein (MAP-1), mRNA
NM_022150	Homo sapiens RFamide-related peptide precursor (RFRP), mRNA
NM_022149	Homo sapiens MAGEF1 protein (MAGEF1), mRNA
NM_022144	Homo sapiens myodulin protein (LOC64102), mRNA
NM_022141	Homo sapiens gamma-parvin (PARVG), mRNA
NM_022134	Homo sapiens glycoprotein beta-Gal 3'-sulfotransferase (GP3ST), mRNA
NM_022131	Homo sapiens calsyntenin-2 (CS2), mRNA
NM_022129	Homo sapiens MAWD binding protein (MAWBP), mRNA
NM_022123	Homo sapiens basic-helix-loop-helix-PAS protein (NPAS3), mRNA
NM_022121	Homo sapiens p53-induced protein PIGPC1 (PIGPC1), mRNA
NM_022120	Homo sapiens hypothetical protein FKSG25 (FLJ00030), mRNA
NM_022114	Homo sapiens PR domain containing 16 (PRDM16), mRNA
NM_022112	Homo sapiens p53-regulated apoptosis-inducing protein 1 (P53AIP1), mRNA
NM_022111	Homo sapiens homolog of Xenopus Claspin (CLASPIN), mRNA
NM_022101	Homo sapiens hypothetical protein FLJ22965 (FLJ22965), mRNA
NM_022087	Homo sapiens hypothetical protein FLJ21634 (FLJ21634), mRNA
NM_022083	Homo sapiens niban protein (NIBAN), mRNA
NM_022078	Homo sapiens hypothetical protein FLJ12455 (FLJ12455), mRNA
NM_022076	Homo sapiens hypothetical protein IMAGE 109914 (LOC63904), mRNA
NM 022072	Homo sapiens hypothetical protein FLJ22609 (FLJ22609), mRNA
NM 022067	Homo sapiens hypothetical protein FLJ12707 (FLJ12707), mRNA
NM 022049	Homo sapiens G-protein coupled receptor 88 (GPR88), mRNA
NM 022044	Homo sapiens stromal cell-derived factor 2-like 1 (SDF2L1), mRNA
NM_022042	Homo sapiens solute carrier family 26 (sulfate transporter), member 1
1111_022072	(SLC26A1), mRNA
NM 022039	Homo sapiens split hand/foot malformation (ectrodactyly) type 3 (SHFM3),
14141_022037	mRNA
NM 021173	Homo sapiens polymerase (DNA-directed), delta 4 (POLD4), mRNA
NM_016371	Homo sapiens hydroxysteroid (17-beta) dehydrogenase 7 (HSD17B7), mRNA
11177 0103/1	1 ATOMIC CAPITORS HYDRONYSICIOID (17-DERS) DEHYDROGERIASE / (MSD1/D/), MRNA

T	1 25000 1 1 1 1 1 1 1 1 1 1 1 1 1 1 1 1 1 1
NM_000023	Homo sapiens sarcoglycan, alpha (50kD dystrophin-associated glycoprotein) (SGCA), mRNA
NM_005099	Homo sapiens a disintegrin-like and metalloprotease (reprolysin type) with thrombospondin type 1 motif, 4 (ADAMTS4), mRNA
NM 016590	Homo sapiens prostate androgen-regulated transcript 1 (PART1), mRNA
	Homo sapiens nuclear transcription factor Y, gamma (NFYC), mRNA
NM_014223 NM_006166	Homo sapiens nuclear transcription factor Y, beta (NFYB), mRNA
	Homo sapiens nuclear transcription factor 1, beta (NY 13), mkNA Homo sapiens karyopherin alpha 4 (importin alpha 3) (KPNA4), mRNA
NM_002268	Homo sapiens ELK1, member of ETS oncogene family (ELK1), mRNA
NM_005229	
NM_021796	Homo sapiens placenta-specific 1 (PLAC1), mRNA Homo sapiens kallikrein 13 (KLK13), mRNA
NM_015596	Homo sapiens kamkrein 13 (KLK13), mktvA Homo sapiens olfactory receptor, family 1, subfamily E, member 1 (OR1E1),
NM_003553	mRNA
NM_021926	Homo sapiens aristaless-like homeobox 4 (ALX4), mRNA
NM_021957	Homo sapiens glycogen synthase 2 (liver) (GYS2), mRNA
NM_020980	Homo sapiens aquaporin 9 (AQP9), mRNA
NM_001614	Homo sapiens actin, gamma 1 (ACTG1), mRNA
NM_018690	Homo sapiens apolipoprotein B48 receptor (APOB48R), mRNA
NM_005230	Homo sapiens ELK3, ETS-domain protein (SRF accessory protein 2) (ELK3), mRNA
NM_003816	Homo sapiens a disintegrin and metalloproteinase domain 9 (meltrin gamma) (ADAM9), mRNA
NM_000847	Homo sapiens glutathione S-transferase A3 (GSTA3), mRNA
NM_021814	Homo sapiens homolog of yeast long chain polyunsaturated fatty acid elongation enzyme 2 (HELO1), mRNA
NM 021628	Homo sapiens arachidonate lipoxygenase 3 (ALOXE3), mRNA
NM 012419	Homo sapiens regulator of G-protein signalling 17 (RGS17), mRNA
NM_014685	Homo sapiens homocysteine-inducible, endoplasmic reticulum stress-inducible, ubiquitin-like domain member 1 (HERPUD1), mRNA
NM 005705	Homo sapiens pan-hematopoietic expression (PHEMX), mRNA
NM 004906	Homo sapiens Wilms' tumour 1-associating protein (KIAA0105), mRNA
NM_003101	Homo sapiens sterol O-acyltransferase (acyl-Coenzyme A cholesterol acyltransferase) 1 (SOAT1), mRNA
NM 021965	Homo sapiens phosphoglucomutase 5 (PGM5), mRNA
NM_003555	Homo sapiens olfactory receptor, family 1, subfamily G, member 1 (OR1G1), mRNA
NM_003552	Homo sapiens olfactory receptor, family 1, subfamily D, member 4 (OR1D4), mRNA
NM 001345	Homo sapiens diacylglycerol kinase, alpha (80kD) (DGKA), mRNA
NM 021620	Homo sapiens PR domain containing 13 (PRDM13), mRNA
NM 020999	Homo sapiens neurogenin 3 (NEUROG3), mRNA
NM 020227	Homo sapiens PR domain containing 9 (PRDM9), mRNA
NM 020226	Homo sapiens PR domain containing 8 (PRDM8), mRNA
NM 020229	Homo sapiens PR domain containing 11 (PRDM11), mRNA
NM 020228	Homo sapiens PR domain containing 10 (PRDM10), mRNA
NM 016412	Homo sapiens insulin-like growth factor 2, antisense (IGF2AS), mRNA
NM 006161	Homo sapiens neurogenin 1 (NEUROG1), mRNA
NM_005734	Homo sapiens homeodomain-interacting protein kinase 3 (HIPK3), mRNA
NM_001818	Homo sapiens aldo-keto reductase family 1, member C4 (chlordecone reductase;
	3-alpha hydroxysteroid dehydrogenase, type I; dihydrodiol dehydrogenase 4) (AKR1C4), mRNA
NM 004363	Homo sapiens carcinoembryonic antigen-related cell adhesion molecule 5

ſ	(CEACAM5), mRNA
NM 002841	Homo sapiens protein tyrosine phosphatase, receptor type, G (PTPRG), mRNA
NM 002716	Homo sapiens protein phosphatase 2 (formerly 2A), regulatory subunit A (PR
1,1,1,_002,10	65), beta isoform (PPP2R1B), mRNA
NM_001785	Homo sapiens cytidine deaminase (CDA), mRNA
NM_003554	Homo sapiens olfactory receptor, family 1, subfamily E, member 2 (OR1E2), mRNA
NM_021961	Homo sapiens TEA domain family member 1 (SV40 transcriptional enhancer factor) (TEAD1), mRNA
NM_002847	Homo sapiens protein tyrosine phosphatase, receptor type, N polypeptide 2 (PTPRN2), mRNA
NM_002778	Homo sapiens prosaposin (variant Gaucher disease and variant metachromatic leukodystrophy) (PSAP), mRNA
NM_000934	Homo sapiens serine (or cysteine) proteinase inhibitor, clade F (alpha-2 antiplasmin, pigment epithelium derived factor), member 2 (SERPINF2), mRNA
NM_000932	Homo sapiens phospholipase C, beta 3 (phosphatidylinositol-specific) (PLCB3), mRNA
NM_000709	Homo sapiens branched chain keto acid dehydrogenase E1, alpha polypeptide (maple syrup urine disease) (BCKDHA), mRNA
NM_001666	Homo sapiens Rho GTPase activating protein 4 (ARHGAP4), mRNA
NM_021815	Homo sapiens solute carrier family 5 (choline transporter), member 7 (SLC5A7), mRNA
NM_014885	Homo sapiens anaphase-promoting complex 10 (APC10), mRNA
NM_021948	Homo sapiens chondroitin sulfate proteoglycan BEHAB/brevican (BCAN), mRNA
NM 021946	Homo sapiens hypothetical protein FLJ11362 (FLJ11362), mRNA
NM_021942	Homo sapiens hypothetical protein FLJ12716 (FLJ12716), mRNA
NM_021940	Homo sapiens hypothetical protein FLJ13159 (FLJ13159), mRNA
NM_021922	Homo sapiens Fanconi anemia, complementation group E (FANCE), mRNA
NM_002644	Homo sapiens polymeric immunoglobulin receptor (PIGR), mRNA
NM_002470	Homo sapiens myosin, heavy polypeptide 3, skeletal muscle, embryonic (MYH3), mRNA
NM_001700	Homo sapiens azurocidin 1 (cationic antimicrobial protein 37) (AZU1), mRNA
NM_003949	Homo sapiens huntingtin-associated protein 1 (neuroan 1) (HAP1), mRNA
NM_021021	Homo sapiens syntrophin, beta 1 (dystrophin-associated protein A1, 59kD, basic component 1) (SNTB1), mRNA
NM_018953	Homo sapiens homeo box C5 (HOXC5), mRNA
NM_012120	Homo sapiens CD2-associated protein (CD2AP), mRNA
NM_007121	Homo sapiens nuclear receptor subfamily 1, group H, member 2 (NR1H2), mRNA
NM_006753	Homo sapiens surfeit 6 (SURF6), mRNA
NM_006200	Homo sapiens proprotein convertase subtilisin/kexin type 5 (PCSK5), mRNA
NM_006426	Homo sapiens dihydropyrimidinase-like 4 (DPYSL4), mRNA
NM_005670	Homo sapiens epilepsy, progressive myoclonus type 2, Lafora disease (laforin) (EPM2A), mRNA
NM_006877	Homo sapiens guanosine monophosphate reductase (GMPR), mRNA
NM_004619	Homo sapiens TNF receptor-associated factor 5 (TRAF5), mRNA
NM_002627	Homo sapiens phosphofructokinase, platelet (PFKP), mRNA
NM_002433	Homo sapiens myelin oligodendrocyte glycoprotein (MOG), mRNA
NM_002207	Homo sapiens integin, alpha 9 (ITGA9), mRNA
NM_002113	Homo sapiens H factor (complement)-like 1 (HFL1), mRNA
NM_002074	Homo sapiens guanine nucleotide binding protein (G protein), beta polypeptide 1

(GNB1), mRNA NM_003733 Homo sapiens 2'-5'oligoadenylate synthetase-like (OASL), mRNA NM_002551 Homo sapiens olfactory receptor, family 3, subfamily A, member 2 (OR3A)	
I NM UUZSSI I Homo sabiens oliactory receptor, family 3, subfamily A, member 2 (ORSA	12)
mRNA	
NM_002389 Homo sapiens membrane cofactor protein (CD46, trophoblast-lymphocyte reactive antigen) (MCP), mRNA	cross-
NM 000870 Homo sapiens 5-hydroxytryptamine (serotonin) receptor 4 (HTR4), mRNA	Δ
NM 000613 Homo sapiens hemopexin (HPX), mRNA	<u>, </u>
NM 000377 Homo sapiens Wiskott-Aldrich syndrome (eczema-thrombocytopenia) (W	(2 A
mRNA	
NM_006981 Homo sapiens nuclear receptor subfamily 4, group A, member 3 (NR4A3) mRNA	' >
NM_000368 Homo sapiens TSC1 gene (hamartin) (TSC1), mRNA	
NM_017416 Homo sapiens interleukin 1 receptor accessory protein-like 2 (IL1RAPL2) mRNA	,
NM_003286 Homo sapiens topoisomerase (DNA) I (TOP1), mRNA	
NM_001068 Homo sapiens topoisomerase (DNA) II beta (180kD) (TOP2B), mRNA	
NM_020470 Homo sapiens putative transmembrane protein; homolog of yeast Golgi	
membrane protein Yiflp (Yiplp-interacting factor) (54TM), mRNA	
NM_006562 Homo sapiens transcription factor similar to D. melanogaster homeodoma:	in
protein lady bird late (LBX1), mRNA	
NM_017545 Homo sapiens hydroxyacid oxidase (glycolate oxidase) 1 (HAO1), mRNA	
NM_002925 Homo sapiens regulator of G-protein signalling 10 (RGS10), mRNA	•
NM_012263 Homo sapiens tubulin tyrosine ligase-like 1 (TTLL1), mRNA	
NM_001212 Homo sapiens complement component 1, q subcomponent binding protein	···
(C1QBP), nuclear gene encoding mitochondrial protein, mRNA	
NM_000491 Homo sapiens complement component 1, q subcomponent, beta polypeptic (C1QB), mRNA	
NM_004720 Homo sapiens endothelial differentiation, lysophosphatidic acid G-protein coupled receptor, 4 (EDG4), mRNA	-
NM_006217 Homo sapiens serine (or cysteine) proteinase inhibitor, clade I (neuroserpi member 2 (SERPINI2), mRNA	n),
NM_018723 Homo sapiens ataxin 2-binding protein 1 (A2BP1), mRNA	
NM_004543 Homo sapiens nebulin (NEB), mRNA	
NM_016151 Homo sapiens prostate derived STE20-like kinase PSK (PSK), mRNA	_
NM_016528 Homo sapiens hydroxyacid oxidase 3 (medium-chain) (HAO3), mRNA	
NM_000185 Homo sapiens serine (or cysteine) proteinase inhibitor, clade D (heparin cofactor), member 1 (SERPIND1), mRNA	
NM_005410 Homo sapiens selenoprotein P, plasma, 1 (SEPP1), mRNA	
NM_005226 Homo sapiens endothelial differentiation, sphingolipid G-protein-coupled	
receptor, 3 (EDG3), mRNA	
NM_005172 Homo sapiens atonal homolog 1 (Drosophila) (ATOH1), mRNA	
NM_005109 Homo sapiens oxidative-stress responsive 1 (OSR1), mRNA	
NM_001498 Homo sapiens glutamate-cysteine ligase, catalytic subunit (GCLC), mRNA	
NM_003922 Homo sapiens heet (homologous to the E6-AP (UBE3A) carboxyl terminu	
domain and RCC1 (CHC1)-like domain (RLD) 1 (HERC1), mRNA	
NM_002061 Homo sapiens glutamate-cysteine ligase, modifier subunit (GCLM), mRN.	A
NM_001088 Homo sapiens arylalkylamine N-acetyltransferase (AANAT), mRNA	
NM 021828 Homo sapiens heparanase-like protein (HPA2), mRNA	
NM_021826 Homo sapiens hypothetical protein FLJ13149 (FLJ13149), mRNA	
NM_021823 Homo sapiens hypothetical protein MDS018 (MDS018), mRNA	
NM_021820 Homo sapiens MDS024 protein (MDS024), mRNA	

NB (021910	Home conions EDCI protein (EDCI) DMA
NM_021819	Homo sapiens ERGL protein (ERGL), mRNA
NM_021818	Homo sapiens WW Domain-Containing Gene (WW45), mRNA
NM_021812	Homo sapiens blepharophimosis, epicanthus inversus and ptosis, candidate 1
ND (021900	(BPESC1), mRNA Homo sapiens TGF(beta)-induced transcription factor 2 (TGIF2), mRNA
NM_021809	
NM_021805	Homo sapiens single Ig IL-1R-related molecule (SIGIRR), mRNA
NM_021803	Homo sapiens interleukin 21 (IL21), mRNA
NM_021798	Homo sapiens interleukin 21 receptor (IL21R), mRNA
NM_020982	Homo sapiens claudin 9 (CLDN9), mRNA
NM_006657	Homo sapiens formiminotransferase cyclodeaminase (FTCD), mRNA
NM_021784	Homo sapiens hepatocyte nuclear factor 3, beta (HNF3B), mRNA
NM_014375	Homo sapiens fetuin B (FETUB), mRNA
NM_021032	Homo sapiens fibroblast growth factor 12 (FGF12), mRNA
NM_019595	Homo sapiens intersectin 2 (ITSN2), mRNA
NM_018991	Homo sapiens DKFZp434A0131 protein (DKFZP434A0131), mRNA
NM_014574	Homo sapiens nuclear autoantigen (GS2NA), mRNA
NM_021002	Homo sapiens interferon, alpha 6 (IFNA6), mRNA
NM_001676	Homo sapiens ATPase, H+/K+ transporting, nongastric, alpha polypeptide (ATP12A), mRNA
NM_019886	Homo sapiens carbohydrate (N-acetylglucosamine 6-O) sulfotransferase 7 (CHST7), mRNA
NM_017581	Homo sapiens cholinergic receptor, nicotinic, alpha polypeptide 9 (CHRNA9), mRNA
NM_001695	Homo sapiens ATPase, H+ transporting, lysosomal (vacuolar proton pump)
	42kD (ATP6C), mRNA
NM 006303	Homo sapiens JTV1 gene (JTV1), mRNA
NM 014413	Homo sapiens heme-regulated initiation factor 2-alpha kinase (HRI), mRNA
NM 012149	Homo sapiens double homeobox, 5 (DUX5), mRNA
NM 012146	Homo sapiens double homeobox, 1 (DUX1), mRNA
NM 021733	Homo sapiens testis-specific kinase substrate (TSKS), mRNA
NM_004339	Homo sapiens pituitary tumor-transforming 1 interacting protein (PTTG1IP), mRNA
NM_004219	Homo sapiens pituitary tumor-transforming 1 (PTTG1), mRNA
NM 003860	Homo sapiens Breakpoint cluster region protein, uterine leiomyoma, 1; barrier to
	autointegration factor (BCRP1), mRNA
NM 007281	Homo sapiens scrapie responsive protein 1 (SCRG1), mRNA
NM 006618	Homo sapiens putative DNA/chromatin binding motif (PLU-1), mRNA
NM 005797	Homo sapiens epithelial V-like antigen 1 (EVA1), mRNA
NM 005508	Homo sapiens chemokine (C-C motif) receptor 4 (CCR4), mRNA
NM 005283	Homo sapiens chemokine (C motif) XC receptor 1 (CCXCR1), mRNA
NM 002547	Homo sapiens oligophrenin 1 (OPHN1), mRNA
NM 020056	Homo sapiens major histocompatibility complex, class II, DQ alpha 2 (HLA-
	DQA2), mRNA
NM_001085	Homo sapiens serine (or cysteine) proteinase inhibitor, clade A (alpha-1
	antiproteinase, antitrypsin), member 3 (SERPINA3), mRNA
NM_013974	Homo sapiens dimethylarginine dimethylaminohydrolase 2 (DDAH2), mRNA
NM_001756	Homo sapiens serine (or cysteine) proteinase inhibitor, clade A (alpha-1
	antiproteinase, antitrypsin), member 6 (SERPINA6), mRNA
NM_000450	Homo sapiens selectin E (endothelial adhesion molecule 1) (SELE), mRNA
NM_006228	Homo sapiens prepronociceptin (PNOC), mRNA
NM_001319	Homo sapiens casein kinase 1, gamma 2 (CSNK1G2), mRNA
NM_000444	Homo sapiens phosphate regulating gene with homologies to endopeptidases on
	The second secon

mRNA NM_021183 Homo sapiens hypothetical protein similar to small G proteins, especia 2A (LOC57826), mRNA NM_021179 Homo sapiens hypothetical protein LOC57821 (LOC57821), mRNA NM_002744 Homo sapiens protein kinase C, zeta (PRKCZ), mRNA NM_000624 Homo sapiens serine (or cysteine) proteinase inhibitor, clade A (alpha-antiproteinase, antitrypsin), member 5 (SERPINA5), mRNA NM_000602 Homo sapiens serine (or cysteine) proteinase inhibitor, clade E (nexin, plasminogen activator inhibitor type 1), member 1 (SERPINE1), mRN NM_020422 Homo sapiens hypothetical protein from clone 24796 (LOC57146), mI	lly RAP-
NM 021179 Homo sapiens hypothetical protein LOC57821 (LOC57821), mRNA NM 002744 Homo sapiens protein kinase C, zeta (PRKCZ), mRNA NM 000624 Homo sapiens serine (or cysteine) proteinase inhibitor, clade A (alpha-antiproteinase, antitrypsin), member 5 (SERPINA5), mRNA NM 000602 Homo sapiens serine (or cysteine) proteinase inhibitor, clade E (nexin, plasminogen activator inhibitor type 1), member 1 (SERPINE1), mRN NM 020422 Homo sapiens hypothetical protein from clone 24796 (LOC57146), mI	
NM 002744 Homo sapiens protein kinase C, zeta (PRKCZ), mRNA NM_000624 Homo sapiens serine (or cysteine) proteinase inhibitor, clade A (alpha-antiproteinase, antitrypsin), member 5 (SERPINA5), mRNA NM_000602 Homo sapiens serine (or cysteine) proteinase inhibitor, clade E (nexin, plasminogen activator inhibitor type 1), member 1 (SERPINE1), mRN NM_020422 Homo sapiens hypothetical protein from clone 24796 (LOC57146), mI	
NM_000624 Homo sapiens serine (or cysteine) proteinase inhibitor, clade A (alpha-antiproteinase, antitrypsin), member 5 (SERPINA5), mRNA NM_000602 Homo sapiens serine (or cysteine) proteinase inhibitor, clade E (nexin, plasminogen activator inhibitor type 1), member 1 (SERPINE1), mRN NM_020422 Homo sapiens hypothetical protein from clone 24796 (LOC57146), mI	
NM_000602 Homo sapiens serine (or cysteine) proteinase inhibitor, clade E (nexin, plasminogen activator inhibitor type 1), member 1 (SERPINE1), mRN NM_020422 Homo sapiens hypothetical protein from clone 24796 (LOC57146), mI	-1
NM_020422 Homo sapiens hypothetical protein from clone 24796 (LOC57146), ml	
NM 020183 Homo sapiens transcription factor BMAL2 (LOC56938), mRNA	
NM 019598 Homo sapiens kallikrein 12 (KLK12), mRNA	
NM 019103 Homo sapiens hypothetical protein (LOC55954), mRNA	
NM_012397 Homo sapiens serine (or cysteine) proteinase inhibitor, clade B (ovalbumember 13 (SERPINB13), mRNA	umin),
NM_000527 Homo sapiens low density lipoprotein receptor (familial hypercholeste (LDLR), mRNA	rolemia)
NM_016200 Homo sapiens U6 snRNA-associated Sm-like protein LSm8 (LOC516) mRNA	91),
NM_014766 Homo sapiens KIAA0193 gene product (KIAA0193), mRNA	
NM 014309 Homo sapiens RNA binding motif protein 9 (RBM9), mRNA	
NM 014080 Homo sapiens dual oxidase-like domains 2 (DUOX2), mRNA	-
NM 014516 Homo sapiens CCR4-NOT transcription complex, subunit 3 (CNOT3)	, mRNA
NM 015032 Homo sapiens KIAA0979 protein (KIAA0979), mRNA	
NM 014656 Homo sapiens KIAA0040 gene product (KIAA0040), mRNA	
NM 015383 Homo sapiens hypothetical protein (DJ328E19.C1.1), mRNA	
NM 012407 Homo sapiens protein kinase C, alpha binding protein (PRKCABP), m	RNA
NM_002208 Homo sapiens integrin, alpha E (antigen CD103, human mucosal lympantigen 1; alpha polypeptide) (ITGAE), mRNA	
NM_002309 Homo sapiens leukemia inhibitory factor (cholinergic differentiation fa	actor)
NM_006919 Homo sapiens serine (or cysteine) proteinase inhibitor, clade B (ovalbumember 3 (SERPINB3), mRNA	umin),
NM_006220 Homo sapiens serine (or cysteine) proteinase inhibitor, clade A (alpha-antiproteinase, antitrypsin), member 2 (SERPINA2), mRNA	-1
NM_006215 Homo sapiens serine (or cysteine) proteinase inhibitor, clade A (alpha-antiproteinase, antitrypsin), member 4 (SERPINA4), mRNA	-1
NM_006021 Homo sapiens deleted in lymphocytic leukemia, 2 (DLEU2), mRNA	
NM_005887 Homo sapiens deleted in lymphocytic leukemia, 1 (DLEU1), mRNA	
NM_005603 Homo sapiens ATPase, Class I, type 8B, member 1 (ATP8B1), mRNA	1
NM 005232 Homo sapiens EphA1 (EPHA1), mRNA	, , , , , , , , , , , , , , , , , , ,
NM_005024 Homo sapiens serine (or cysteine) proteinase inhibitor, clade B (ovalbumember 10 (SERPINB10), mRNA	umin),
NM_004779 Homo sapiens CCR4-NOT transcription complex, subunit 8 (CNOT8)	, mRNA
NM_004155 Homo sapiens serine (or cysteine) proteinase inhibitor, clade B (ovalbe member 9 (SERPINB9), mRNA	
NM_004568 Homo sapiens serine (or cysteine) proteinase inhibitor, clade B (ovalbe member 6 (SERPINB6), mRNA	umin),
NM_004408 Homo sapiens dynamin 1 (DNM1), mRNA	
NM_004409 Homo sapiens dystrophia myotonica-protein kinase (DMPK), mRNA	
NM_004717 Homo sapiens diacylglycerol kinase, iota (DGKI), mRNA	

NM_000214	Homo sapiens jagged 1 (Alagille syndrome) (JAG1), mRNA
NM_001347	Homo sapiens diacylglycerol kinase, theta (110kD) (DGKQ), mRNA
NM_003454	Homo sapiens zinc finger protein 200 (ZNF200), mRNA
NM_003334	Homo sapiens ubiquitin-activating enzyme E1 (A1S9T and BN75 temperature sensitivity complementing) (UBE1), mRNA
NM_000354	Homo sapiens serine (or cysteine) proteinase inhibitor, clade A (alpha-1
14141_000554	antiproteinase, antitrypsin), member 7 (SERPINA7), mRNA
NM_000945	Homo sapiens protein phosphatase 3 (formerly 2B), regulatory subunit B (19kD),
_	alpha isoform (calcineurin B, type I) (PPP3R1), mRNA
NM 000305	Homo sapiens paraoxonase 2 (PON2), mRNA
NM 000928	Homo sapiens phospholipase A2, group IB (pancreas) (PLA2G1B), nuclear gene
_	encoding mitochondrial protein, mRNA
NM_000295	Homo sapiens serine (or cysteine) proteinase inhibitor, clade A (alpha-1
_	antiproteinase, antitrypsin), member 1 (SERPINA1), mRNA
NM_002640	Homo sapiens serine (or cysteine) proteinase inhibitor, clade B (ovalbumin),
_	member 8 (SERPINB8), mRNA
NM_002639	Homo sapiens serine (or cysteine) proteinase inhibitor, clade B (ovalbumin),
_	member 5 (SERPINB5), mRNA
NM_002615	Homo sapiens serine (or cysteine) proteinase inhibitor, clade F (alpha-2
_	antiplasmin, pigment epithelium derived factor), member 1 (SERPINF1), mRNA
NM_002575	Homo sapiens serine (or cysteine) proteinase inhibitor, clade B (ovalbumin),
	member 2 (SERPINB2), mRNA
NM_000220	Homo sapiens potassium inwardly-rectifying channel, subfamily J, member 1
	(KCNJ1), mRNA
NM_000191	Homo sapiens 3-hydroxymethyl-3-methylglutaryl-Coenzyme A lyase
	(hydroxymethylglutaricaciduria) (HMGCL), mRNA
NM_001978	Homo sapiens erythrocyte membrane protein band 4.9 (dematin) (EPB49),
	mRNA
NM_003646	Homo sapiens diacylglycerol kinase, zeta (104kD) (DGKZ), mRNA
NM_001346	Homo sapiens diacylglycerol kinase, gamma (90kD) (DGKG), mRNA
NM_003647	Homo sapiens diacylglycerol kinase, epsilon (64kD) (DGKE), mRNA
NM_001235	Homo sapiens serine (or cysteine) proteinase inhibitor, clade H (heat shock protein 47), member 2 (SERPINH2), mRNA
NM 001694	Homo sapiens ATPase, H+ transporting, lysosomal (vacuolar proton pump)
	16kD (ATP6L), mRNA
NM 000488	Homo sapiens serine (or cysteine) proteinase inhibitor, clade C (antithrombin),
	member 1 (SERPINC1), mRNA
NM 021156	Homo sapiens hypothetical protein (DJ971N18.2), mRNA
NM 000875	Homo sapiens insulin-like growth factor 1 receptor (IGF1R), mRNA
NM_000605	Homo sapiens interferon, alpha 2 (IFNA2), mRNA
NM 021647	Homo sapiens KIAA0626 gene product (KIAA0626), mRNA
NM 021645	Homo sapiens KIAA0266 gene product (KIAA0266), mRNA
NM_021049	Homo sapiens thymosin, beta 4, X chromosome (TMSB4X), mRNA
NM 021642	Homo sapiens for fragment of IgG, low affinity IIa, receptor for (CD32)
1111_021042	(FCGR2A), mRNA
NM 021240	Homo sapiens testis-specific protein (LOC58524), mRNA
NM 021189	Homo sapiens hypothetical protein FLJ10698 (LOC57863), mRNA
NM 021129	Homo sapiens pyrophosphatase (inorganic) (PP), nuclear gene encoding
11111_021129	mitochondrial protein, mRNA
NM 015140	Homo sapiens KIAA0153 protein (KIAA0153), mRNA
NM 021635	Homo sapiens UC28 protein (UC28), mRNA
NM 021631	Homo sapiens OC28 protein (OC28), inRNA Homo sapiens apoptosis inhibitor (FKSG2), mRNA
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NM_021615	Homo sapiens carbohydrate (N-acetylglucosamine 6-0) sulfotransferase 6 (CHST6), mRNA
NM 012334	Homo sapiens myosin X (MYO10), mRNA
NM 020363	Homo sapiens deleted in azoospermia 2 (DAZ2), mRNA
NM 020364	Homo sapiens deleted in azoospermia 3 (DAZ3), mRNA
NM 017445	Homo sapiens H2B histone family, member S (H2BFS), mRNA
NM_021132	Homo sapiens protein phosphatase 3 (formerly 2B), catalytic subunit, beta
	isoform (calcineurin A beta) (PPP3CB), mRNA
NM_021016	Homo sapiens pregnancy specific beta-1-glycoprotein 3 (PSG3), mRNA
NM_015705	Homo sapiens hypothetical protein (DJ1042K10.2), mRNA
NM_021572	Homo sapiens ectonucleotide pyrophosphatase/phosphodiesterase 5 (putative function) (ENPP5), mRNA
NM_021216	Homo sapiens endothelial zinc finger protein induced by tumor necrosis factor
_	alpha (EZFIT), mRNA
NM 001332	Homo sapiens catenin (cadherin-associated protein), delta 2 (neural plakophilin-
_	related arm-repeat protein) (CTNND2), mRNA
NM_021185	Homo sapiens hypothetical protein DKFZp434A1022 (DKFZP434A1022),
	mRNA
NM_018955	Homo sapiens ubiquitin B (UBB), mRNA
NM_017533	Homo sapiens myosin, heavy polypeptide 4, skeletal muscle (MYH4), mRNA
NM_014621	Homo sapiens homeo box D4 (HOXD4), mRNA
NM_000618	Homo sapiens insulin-like growth factor 1 (somatomedia C) (IGF1), mRNA
NM_021571	Homo sapiens ICEBERG caspase-1 inhibitor (ICEBERG), mRNA
NM_000045	Homo sapiens arginase, liver (ARG1), mRNA
NM_005692	Homo sapiens ATP-binding cassette, sub-family F (GCN20), member 2 (ABCF2), mRNA
NB (001000	
NM_001090	Homo sapiens ATP-binding cassette, sub-family F (GCN20), member 1 (ABCF1), mRNA
NM_002858	Homo sapiens ATP-binding cassette, sub-family D (ALD), member 3 (ABCD3), mRNA
NM_001172	Homo sapiens arginase, type II (ARG2), nuclear gene encoding mitochondrial protein, mRNA
NM_001117	Homo sapiens adenylate cyclase activating polypeptide 1 (pituitary)
	(ADCYAP1), mRNA
NM 004036	Homo sapiens adenylate cyclase 3 (ADCY3), mRNA
NM 019843	Homo sapiens eIF4E-transporter (4E-T), mRNA
NM_006454	Homo sapiens Mad4 homolog (MAD4), mRNA
NM_002355	Homo sapiens mannose-6-phosphate receptor (cation dependent) (M6PR), mRNA
NM 014287	Homo sapiens pM5 protein (PM5), mRNA
NM 004102	Homo sapiens fatty acid binding protein 3, muscle and heart (mammary-derived
	growth inhibitor) (FABP3), mRNA
NM_000134	Homo sapiens fatty acid binding protein 2, intestinal (FABP2), mRNA
NM_005354	Homo sapiens jun D proto-oncogene (JUND), mRNA
NM_005159	Homo sapiens actin, alpha, cardiac muscle (ACTC), mRNA
NM_019848	Homo sapiens Protein P3 (P3), mRNA
NM_003948	Homo sapiens cyclin-dependent kinase-like 2 (CDC2-related kinase) (CDKL2), mRNA
NM_021131	Homo sapiens protein phosphatase 2A, regulatory subunit B' (PR 53) (PPP2R4), mRNA
NM_021268	Homo sapiens interferon, alpha 17 (IFNA17), mRNA
NM 002339	Homo sapiens lymphocyte-specific protein 1 (LSP1), mRNA

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NM_001166	Homo sapiens baculoviral IAP repeat-containing 2 (BIRC2), mRNA
NM_003399	Homo sapiens X-prolyl aminopeptidase (aminopeptidase P) 2, membrane-bound
) T. f. 000541	(XPNPEP2), mRNA
NM_000541	Homo sapiens S-antigen; retina and pineal gland (arrestin) (SAG), mRNA
NM_013262	Homo sapiens myosin regulatory light chain interacting protein (MIR), mRNA
NM_005393	Homo sapiens plexin B3 (PLXNB3), mRNA
NM_021098	Homo sapiens calcium channel, voltage-dependent, alpha 1H subunit
	(CACNA1H), mRNA
NM_021257	Homo sapiens neuroglobin (NGB), mRNA
NM_021253	Homo sapiens ring finger protein 23 (RNF23), mRNA
NM_021247	Homo sapiens protamine 3 (PRM3), mRNA
NM_021242	Homo sapiens hypothetical protein STRAIT11499 (STRAIT11499), mRNA
NM_021238	Homo sapiens TERA protein (TERA), mRNA
NM_021223	Homo sapiens myosin light chain 2a (LOC58498), mRNA
NM_021221	Homo sapiens G5b protein (G5B), mRNA
NM_021210	Homo sapiens MUM2 protein (MUM2), mRNA
NM_021208	Homo sapiens EST-YD1 protein (EST-YD1), mRNA
NM_021200	Homo sapiens PH domain containing protein in retina 1 (PHRET1), mRNA
NM_021199	Homo sapiens CGI-44 protein; sulfide dehydrogenase like (yeast) (CGI-44),
	mRNA
NM_021198	Homo sapiens nuclear LIM interactor-interacting factor (NLI-IF), mRNA
NM_021193	Homo sapiens homeo box D12 (HOXD12), mRNA
NM_021192	Homo sapiens homeo box D11 (HOXD11), mRNA
NM_021188	Homo sapiens clones 23667 and 23775 zinc finger protein (LOC57862), mRNA
NM_021184	Homo sapiens G4 protein (G4), mRNA
NM_021177	Homo sapiens U6 snRNA-associated Sm-like protein (LSM2), mRNA
NM_021174	Homo sapiens p30 DBC protein (LOC57805), mRNA
NM_021167	Homo sapiens hypothetical protein WUGSC:H_RG083M05.2 (LOC57798),
	mRNA
NM_021159	Homo sapiens RAP1, GTP-GDP dissociation stimulator 1 (RAP1GDS1), mRNA
NM_021155	Homo sapiens CD209 antigen (CD209), mRNA
NM_021147	Homo sapiens uracil-DNA glycosylase 2 (UNG2), mRNA
NM_021140	Homo sapiens ubiquitously transcribed tetratricopeptide repeat gene, X
	chromosome (UTX), mRNA
NM_021139	Homo sapiens UDP glycosyltransferase 2 family, polypeptide B4 (UGT2B4),
	mRNA
NM_021138	Homo sapiens TNF receptor-associated factor 2 (TRAF2), mRNA
NM_021137	Homo sapiens tumor necrosis factor, alpha-induced protein 1 (endothelial)
	(TNFAIP1), mRNA
NM_021136	Homo sapiens reticulon 1 (RTN1), mRNA
NM_021135	Homo sapiens ribosomal protein S6 kinase, 90kD, polypeptide 2 (RPS6KA2),
	mRNA
NM_021133	Homo sapiens ribonuclease L (2',5'-oligoisoadenylate synthetase-dependent)
	(RNASEL), mRNA
NM_021130	Homo sapiens peptidylprolyl isomerase A (cyclophilin A) (PPIA), mRNA
NM_021120	Homo sapiens discs, large (Drosophila) homolog 3 (neuroendocrine-dlg)
	(DLG3), mRNA
NM_004239	Homo sapiens thyroid hormone receptor interactor 11 (TRIP11), mRNA
NM_004238	Homo sapiens thyroid hormone receptor interactor 12 (TRIP12), mRNA
NM_004745	Homo sapiens discs, large (Drosophila) homolog-associated protein 2
ND4 004697	(DLGAP2), mRNA
NM_004687	Homo sapiens myotubularin related protein 4 (MTMR4), mRNA

NM 004348	Homo sapiens runt-related transcription factor 2 (RUNX2), mRNA
NM_004348 NM_021096	Homo sapiens calcium channel, voltage-dependent, alpha 1I subunit
14147 051030	(CACNA11), mRNA
NM 021105	Homo sapiens phospholipid scramblase 1 (PLSCR1), mRNA
NM 002957	Homo sapiens retinoid X receptor, alpha (RXRA), mRNA
NM 006268	Homo sapiens requiem, apoptosis response zinc finger gene (REQ), mRNA
NM 001106	Homo sapiens activin A receptor, type IIB (ACVR2B), mRNA
NM 001616	Homo sapiens activin A receptor, type II (ACVR2), mRNA
NM 001105	Homo sapiens activin A receptor, type I (ACVR1), mRNA
NM 005570	Homo sapiens lectin, mannose-binding, 1 (LMAN1), mRNA
NM_021083	Homo sapiens Kell blood group precursor (McLeod phenotype) (XK), mRNA
NM 013258	Homo sapiens apoptosis-associated speck-like protein containing a CARD
11111_013250	(ASC), mRNA
NM 006518	Homo sapiens small proline-rich protein 2C (SPRR2C), mRNA
NM 006507	Homo sapiens regenerating islet-derived 1 beta (pancreatic stone protein,
- 12/200000	pancreatic thread protein) (REG1B), mRNA
NM 006563	Homo sapiens Kruppel-like factor 1 (erythroid) (KLF1), mRNA
NM_006258	Homo sapiens protein kinase, cGMP-dependent, type I (PRKG1), mRNA
NM 006353	Homo sapiens high-mobility group (nonhistone chromosomal) protein 17-like 3
L	(HMG17L3), mRNA
NM_005987	Homo sapiens small proline-rich protein 1A (SPRR1A), mRNA
NM_005952	Homo sapiens metallothionein 1X (MT1X), mRNA
NM_005950	Homo sapiens metallothionein 1G (MT1G), mRNA
NM_005699	Homo sapiens interleukin 18 binding protein (IL18BP), mRNA
NM_004618	Homo sapiens topoisomerase (DNA) III alpha (TOP3A), mRNA
NM_001136	Homo sapiens advanced glycosylation end product-specific receptor (AGER), mRNA
NM_000866	Homo sapiens 5-hydroxytryptamine (serotonin) receptor 1F (HTR1F), mRNA
NM_000637	Homo sapiens glutathione reductase (GSR), mRNA
NM_000636	Homo sapiens superoxide dismutase 2, mitochondrial (SOD2), mRNA
NM_000635	Homo sapiens regulatory factor X, 2 (influences HLA class II expression) (RFX2), mRNA
NM_000629	Homo sapiens interferon (alpha, beta and omega) receptor 1 (IFNAR1), mRNA
NM_000625	Homo sapiens nitric oxide synthase 2A (inducible, hepatocytes) (NOS2A), mRNA
NM_003998	Homo sapiens nuclear factor of kappa light polypeptide gene enhancer in B-cells 1 (p105) (NFKB1), mRNA
NM_000621	Homo sapiens 5-hydroxytryptamine (serotonin) receptor 2A (HTR2A), mRNA
NM_000620	Homo sapiens nitric oxide synthase 1 (neuronal) (NOS1), mRNA
NM_000619	Homo sapiens interferon, gamma (IFNG), mRNA
NM_000617	Homo sapiens solute carrier family 11 (proton-coupled divalent metal ion
NM_000616	transporters), member 2 (SLC11A2), mRNA Homo sapiens CD4 antigen (p55) (CD4), mRNA
NM_000611	Homo sapiens CD59 antigen p18-20 (antigen identified by monoclonal
11111_000011	antibodies 16.3A5, EJ16, EJ30, EL32 and G344) (CD59), mRNA
NM_000610	Homo sapiens CD44 antigen (homing function and Indian blood group system)
	(CD44), mRNA
NM 000603	Homo sapiens nitric oxide synthase 3 (endothelial cell) (NOS3), mRNA
NM_000597	Homo sapiens insulin-like growth factor binding protein 2 (36kD) (IGFBP2),
	mRNA
NM_000594	Homo sapiens tumor necrosis factor (TNF superfamily, member 2) (TNF), mRNA

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NM_000585	Homo sapiens interleukin 15 (IL15), mRNA
NM_000586	Homo sapiens interleukin 2 (IL2), mRNA
NM_000577	Homo sapiens interleukin 1 receptor antagonist (IL1RN), mRNA
NM_000576	Homo sapiens interleukin 1, beta (IL1B), mRNA
NM_000574	Homo sapiens decay accelerating factor for complement (CD55, Cromer blood group system) (DAF), mRNA
NM_000572	Homo sapiens interleukin 10 (IL10), mRNA
NM_000570	Homo sapiens Fc fragment of IgG, low affinity IIIb, receptor for (CD16) (FCGR3B), mRNA
NM_000567	Homo sapiens C-reactive protein, pentraxin-related (CRP), mRNA
NM_000566	Homo sapiens Fc fragment of IgG, high affinity Ia, receptor for (CD64) (FCGR1A), mRNA
NM_000564	Homo sapiens interleukin 5 receptor, alpha (IL5RA), mRNA
NM_000561	Homo sapiens glutathione S-transferase M1 (GSTM1), mRNA
NM_000555	Homo sapiens doublecortex; lissencephaly, X-linked (doublecortin) (DCX), mRNA
NM_000298	Homo sapiens pyruvate kinase, liver and RBC (PKLR), nuclear gene encoding mitochondrial protein, mRNA
NM_000259	Homo sapiens myosin VA (heavy polypeptide 12, myoxin) (MYO5A), mRNA
NM_000525	Homo sapiens potassium inwardly-rectifying channel, subfamily J, member 11 (KCNJ11), mRNA
NM_021090	Homo sapiens myotubularin related protein 3 (MTMR3), mRNA
NM 021077	Homo sapiens neuromedin B (NMB), mRNA
NM 021068	Homo sapiens interferon, alpha 4 (IFNA4), mRNA
NM 006512	Homo sapiens serum amyloid A4, constitutive (SAA4), mRNA
NM 006607	Homo sapiens pituitary tumor-transforming 2 (PTTG2), mRNA
NM_021075	Homo sapiens NADH dehydrogenase (ubiquinone) flavoprotein 3 (10kD) (NDUFV3), mRNA
NM 005951	Homo sapiens metallothionein 1H (MT1H), mRNA
NM_000330	Homo sapiens retinoschisis (X-linked, juvenile) 1 (RS1), mRNA
NM_005597	Homo sapiens nuclear factor I/C (CCAAT-binding transcription factor) (NFIC), mRNA
NM 005268	Homo sapiens gap junction protein, beta 5 (connexin 31.1) (GJB5), mRNA
NM_004268	Homo sapiens cofactor required for Sp1 transcriptional activation, subunit 6 (77kD) (CRSP6), mRNA
NM_004355	Homo sapiens CD74 antigen (invariant polypeptide of major histocompatibility complex, class II antigen-associated) (CD74), mRNA
NM 002760	Homo sapiens protein kinase, Y-linked (PRKY), mRNA
NM_002520	Homo sapiens nucleophosmin (nucleolar phosphoprotein B23, numatrin) (NPM1), mRNA
NM_002167	Homo sapiens inhibitor of DNA binding 3, dominant negative helix-loop-helix protein (ID3), mRNA
NM_002028	Homo sapiens farnesyltransferase, CAAX box, beta (FNTB), mRNA
NM_003491	Homo sapiens N-acetyltransferase, homolog of S. cerevisiae ARD1 (ARD1), mRNA
NM 001770	Homo sapiens CD19 antigen (CD19), mRNA
NM 001664	Homo sapiens ras homolog gene family, member A (ARHA), mRNA
NM 003919	Homo sapiens sarcoglycan, epsilon (SGCE), mRNA
NM_003841	Homo sapiens tumor necrosis factor receptor superfamily, member 10c, decoy without an intracellular domain (TNFRSF10C), mRNA
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NM 003455	Homo sapiens zinc finger protein 202 (ZNF202), mRNA

NM_003166 Homo sapiens tetratricopeptide repeat domain 3 (TTC3), mRNA NM_003166 Homo sapiens sulfotransferase family, cytosolic, 1A, phenol-preferring, member 3 (SULT1A3), mRNA NM_003117 Homo sapiens sperm adhesion molecule 1 (PH-20 hyaluronidase, zona pellucida binding) (SPAM1), mRNA NM_002222 Homo sapiens inositol 1,4,5-triphosphate receptor, type 1 (ITPR1), mRNA NM_001532 Homo sapiens solute carrier family 29 (nucleoside transporters), member 2 (SLC29A2), mRNA NM_001437 Homo sapiens estrogen receptor 2 (ER beta) (ESR2), mRNA NM_001331 Homo sapiens catenin (cadherin-associated protein), delta 1 (CTNND1), mRNA NM_001307 Homo sapiens claudin 7 (CLDN7), mRNA NM_001194 Homo sapiens hyperpolarization activated cyclic nucleotide-gated potassium channel 2 (HCN2), mRNA NM_001175 Homo sapiens Rho GDP dissociation inhibitor (GDI) beta (ARHGDIB), mRNA NM_000641 Homo sapiens interleukin 11 (IL11), mRNA NM_000640 Homo sapiens interleukin 13 receptor, alpha 2 (IL13RA2), mRNA NM_000609 Homo sapiens neural cell adhesion molecule 1 (NCAM1), mRNA NM_000609 Homo sapiens stromal cell-derived factor 1 (SDF1), mRNA NM_000600 Homo sapiens interleukin 6 (interferon, beta 2) (IL6), mRNA NM_000599 Homo sapiens interleukin 9 (IL9), mRNA
NM_003166 Homo sapiens sulfotransferase family, cytosolic, 1A, phenol-preferring, member 3 (SULT1A3), mRNA NM_003117 Homo sapiens sperm adhesion molecule 1 (PH-20 hyaluronidase, zona pellucida binding) (SPAM1), mRNA NM_002222 Homo sapiens inositol 1,4,5-triphosphate receptor, type 1 (ITPR1), mRNA NM_001532 Homo sapiens solute carrier family 29 (nucleoside transporters), member 2 (SLC29A2), mRNA NM_001437 Homo sapiens estrogen receptor 2 (ER beta) (ESR2), mRNA NM_001331 Homo sapiens catenin (cadherin-associated protein), delta 1 (CTNND1), mRNA NM_001307 Homo sapiens claudin 7 (CLDN7), mRNA NM_001194 Homo sapiens hyperpolarization activated cyclic nucleotide-gated potassium channel 2 (HCN2), mRNA NM_001175 Homo sapiens Rho GDP dissociation inhibitor (GDI) beta (ARHGDIB), mRNA NM_00036 Homo sapiens interleukin 11 (IL11), mRNA NM_000641 Homo sapiens interleukin 13 receptor, alpha 2 (IL13RA2), mRNA NM_000640 Homo sapiens neural cell adhesion molecule 1 (NCAM1), mRNA NM_000609 Homo sapiens interleukin 6 (interferon, beta 2) (IL6), mRNA NM_000599 Homo sapiens interleukin 9 (IL9), mRNA NM_000590 Homo sapiens interleukin 9 (IL9), mRNA
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NM_001532 Homo sapiens solute carrier family 29 (nucleoside transporters), member 2 (SLC29A2), mRNA NM_001437 Homo sapiens estrogen receptor 2 (ER beta) (ESR2), mRNA NM_001301 Homo sapiens catenin (cadherin-associated protein), delta 1 (CTNND1), mRNA NM_001307 Homo sapiens claudin 7 (CLDN7), mRNA NM_001194 Homo sapiens hyperpolarization activated cyclic nucleotide-gated potassium channel 2 (HCN2), mRNA NM_001175 Homo sapiens Rho GDP dissociation inhibitor (GDI) beta (ARHGDIB), mRNA NM_000936 Homo sapiens pancreatic lipase (PNLIP), mRNA NM_000641 Homo sapiens interleukin 11 (IL11), mRNA NM_000640 Homo sapiens interleukin 13 receptor, alpha 2 (IL13RA2), mRNA NM_000615 Homo sapiens neural cell adhesion molecule 1 (NCAM1), mRNA NM_000609 Homo sapiens stromal cell-derived factor 1 (SDF1), mRNA NM_000600 Homo sapiens interleukin 6 (interferon, beta 2) (IL6), mRNA NM_000590 Homo sapiens interleukin 9 (IL9), mRNA
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NM_000590 Homo sapiens interleukin 9 (IL9), mRNA
AD C 000504 TY : : 1 11 0 5 5
NM_000584 Homo sapiens interleukin 8 (IL8), mRNA
NM 000581 Homo sapiens glutathione peroxidase 1 (GPX1), mRNA
NM_000560 Homo sapiens CD53 antigen (CD53), mRNA
NM 000528 Homo sapiens mannosidase, alpha, class 2B, member 1 (MAN2B1), mRNA
NM_000404 Homo sapiens galactosidase, beta 1 (GLB1), mRNA
NM_001275 Homo sapiens chromogranin A (parathyroid secretory protein 1) (CHGA), mRNA
NM_006768 Homo sapiens BRCA1 associated protein (BRAP), mRNA
NM_003469 Homo sapiens secretogranin II (chromogranin C) (SCG2), mRNA
NM_012326 Homo sapiens microtubule-associated protein, RP/EB family, member 3
(MAPRE3), mRNA
NM_021057 Homo sapiens interferon, alpha 7 (IFNA7), mRNA
NM_021062 Homo sapiens H2B histone family, member F (H2BFF), mRNA
NM 021063 Homo sapiens H2B histone family, member B (H2BFB), mRNA
NM_021065 Homo sapiens H2A histone family, member G (H2AFG), mRNA
NM_004146 Homo sapiens NADH dehydrogenase (ubiquinone) 1 beta subcomplex, 7 (18kD,
NM_001746 Homo sapiens calnexin (CANX), mRNA
NM_003661 Homo sapiens apolipoprotein L (APOL), mRNA
NM_021052 Homo sapiens H2A histone family, member A (H2AFA), mRNA
NM_020988 Homo sapiens guanine nucleotide binding protein (G protein), alpha activating
activity polypeptide O (GNAO1), mRNA
NM_000133 Homo sapiens coagulation factor IX (plasma thromboplastic component,
Christmas disease, hemophilia B) (F9), mRNA
NM 000130 Homo sapiens coagulation factor V (proaccelerin, labile factor) (F5), mRNA
NM 001993 Homo sapiens coagulation factor III (thromboplastin, tissue factor) (F3), mRNA
NM_020689 Homo sapiens sodium calcium exchanger (NCKX3), mRNA
NM_021033 Homo sapiens RAP2A, member of RAS oncogene family (RAP2A), mRNA
NM 021023 Homo sapiens complement factor H related 3 (FHR-3), mRNA
NM_021026 Homo sapiens ret finger protein-like ! (RFPL1), mRNA

Homo sapiens suppressin (nuclear deformed epidermal autoregulatory factor-1 (DEAF-1)-related) (SPN), mRNA
Homo sapiens B-cell CLL/lymphoma 7A (BCL7A), mRNA
Homo sapiens cancer/testis antigen 2 (CTAG2), mRNA
Homo sapiens pituitary tumor-transforming 3 (PTTG3), mRNA
Homo sapiens left-right determination, factor B (LEFTB), mRNA
Homo sapiens synovial sarcoma, X breakpoint 3 (SSX3), mRNA
Homo sapiens synovial sarcoma, X breakpoint 5 (SSX5), mRNA
Homo sapiens sodium channel, voltage-gated, type II, alpha 2 polypeptide
(SCN2A2), mRNA Homo sapiens potassium inwardly-rectifying channel, subfamily J, member 12
(KCNJ12), mRNA
Homo sapiens haptoglobin-related protein (HPR), mRNA
Homo sapiens spectrin, beta, erythrocytic (includes spherocytosis, clinical type I) (SPTB), mRNA
Homo sapiens putative nuclear protein (HRIHFB2122), mRNA
Homo sapiens casein kinase 2, beta polypeptide (CSNK2B), mRNA
Homo sapiens C-type (calcium dependent, carbohydrate-recognition domain) lectin, superfamily member 5 (CLECSF5), mRNA
Homo sapiens amylase, alpha 2B; pancreatic (AMY2B), mRNA
Homo sapiens zinc finger protein 275 (ZNF275), mRNA
Homo sapiens anti-Mullerian hormone receptor, type II (AMHR2), mRNA
Homo sapiens CEGP1 protein (CEGP1), mRNA
Homo sapiens HT018 protein (HT018), mRNA
Homo sapiens lipase protein (LOC57406), mRNA
Homo sapiens \$100-type calcium binding protein A14 (LOC57402), mRNA
Homo sapiens activation-induced cytidine deaminase (AICDA), mRNA
Homo sapiens zinc finger protein 304 (ZNF304), mRNA
Homo sapiens sentrin/SUMO-specific protease (SENP7), mRNA
Homo sapiens reserved (ASCL3), mRNA
Homo sapiens RP42 homolog (RP42), mRNA
Homo sapiens ankyrin repeat domain 3 (ANKRD3), mRNA
Homo sapiens ATPase, H(+)-transporting, lysosomal, noncatalytic accessory
Homo sapiens Al Pase, ri(+)-transporting, lysosomal, noncatalytic accessory
protein 1B (ATP6N1B), mRNA
Homo sapiens twisted gastrulation (TSG), mRNA
Homo sapiens G protein-coupled receptor 85 (GPR85), mRNA
Homo sapiens sphingosine-1-phosphate lyase 1 (SGPL1), mRNA
Homo sapiens chromobox homolog 6 (CBX6), mRNA
Homo sapiens homeo box A2 (HOXA2), mRNA
Homo sapiens similar to prokaryotic-type class I peptide chain release factors (LOC54516), mRNA
Homo sapiens tight junction protein 3 (zona occludens 3) (TJP3), mRNA
Homo sapiens hypothetical protein dJ122O8.2 (DJ122O8.2), mRNA
Homo sapiens hypothetical protein dJ462O23.2 (DJ462O23.2), mRNA
Homo sapiens hypothetical protein DKFZp586E1923 (DKFZP586E1923), mRNA
Homo sapiens hypothetical protein A-211C6.1 (LOC57149), mRNA
Homo sapiens hypothetical protein dJ465N24.2.1 (DJ465N24.2.1), mRNA
Homo sapiens hypothetical protein dJ37E16.5 (DJ37E16.5), mRNA
Homo sapiens hypothetical protein (LOC57019), mRNA
Homo sapiens olfactory receptor, family 2, subfamily S, member 2 (OR2S2), mRNA

NM 019605	Homo sapiens hypothetical protein (DJ667H12.2), mRNA
NM_019601	Homo sapiens Sushi domain (SCR repeat) containing (BK65A6.2), mRNA
NM 018433	Homo sapiens putative zinc finger protein (LOC55818), mRNA
NM 019095	Homo sapiens hypothetical protein (LOC54675), mRNA
NM_019089	Homo sapiens hairy and enhancer of split (Drosophila) homolog 2 (HES2),
1111_01505	mRNA
NM 018982	Homo sapiens hypothetical protein (DJ167A19.1), mRNA
NM 018974	Homo sapiens unc93 (C.elegans) homolog A (UNC93A), mRNA
NM_014499	Homo sapiens putative purinergic receptor (P2Y10), mRNA
NM 020530	Homo sapiens oncostatin M (OSM), mRNA
NM 020529	Homo sapiens nuclear factor of kappa light polypeptide gene enhancer in B-cells
111.2_0200	inhibitor, alpha (NFKBIA), mRNA
NM 014204	Homo sapiens BCL2-related ovarian killer (BOK), mRNA
NM 020527	Homo sapiens HUG1 gene (HUG1), mRNA
NM 006093	Homo sapiens proteoglycan 3 (PRG3), mRNA
NM 020533	Homo sapiens mucolipin 1 (MCOLN1), mRNA
NM 007345	Homo sapiens zinc finger protein 236 (ZNF236), mRNA
NM 002217	Homo sapiens pre-alpha (globulin) inhibitor, H3 polypeptide (ITIH3), mRNA
NM 018693	Homo sapiens vitiligo-associated protein VIT-1 (VIT1), mRNA
NM 006777	Homo sapiens Kaiso (ZNF-kaiso), mRNA
NM 020436	Homo sapiens similar to SALL1 (sal (Drosophila)-like (LOC57167), mRNA
NM 020142	Homo sapiens NADH:ubiquinone oxidoreductase MLRQ subunit homolog
-	(LOC56901), mRNA
NM 020123	Homo sapiens endomembrane protein emp70 precursor isolog (LOC56889),
	mRNA.
NM_018845	Homo sapiens stromal cell protein (LOC55974), mRNA
NM_018842	Homo sapiens insulin receptor tyrosine kinase substrate (LOC55971), mRNA
NM_018841	Homo sapiens G-protein gamma-12 subunit (LOC55970), mRNA
NM_018839	Homo sapiens p47 protein (LOC55968), mRNA
NM_016352	Homo sapiens carboxypeptidase A3 (LOC51200), mRNA
NM_016302	Homo sapiens protein x 0001 (LOC51185), mRNA
NM_014332	Homo sapiens small muscle protein, X-linked (SMPX), mRNA
NM_018948	Homo sapiens Gene 33/Mig-6 (MIG-6), mRNA
NM_014587	Homo sapiens SRY (sex determining region Y)-box 8 (SOX8), mRNA
NM_005745	Homo sapiens accessory proteins BAP31/BAP29 (DXS1357E), mRNA
NM_001094	Homo sapiens amiloride-sensitive cation channel 1, neuronal (degenerin)
	(ACCN1), mRNA
NM_019609	Homo sapiens metallocarboxypeptidase CPX-1 (CPX-1), mRNA
NM_018844	Homo sapiens B-cell receptor-associated protein BAP29 (BAP29), mRNA
NM_017572	Homo sapiens G protein-coupled receptor kinase 7 (GPRK7), mRNA
NM_016418	Homo sapiens clone FLB5214 (LOC51219), mRNA
NM_016301	Homo sapiens protein x 0004 (LOC51184), mRNA
NM_013387	Homo sapiens ubiquinol-cytochrome c reductase complex (7.2 kD) (HSPC051),
2000000	mRNA
NM_020469	Homo sapiens ABO blood group (transferase A, alpha 1-3-N-
	acetylgalactosaminyltransferase; transferase B, alpha 1-3-galactosyltransferase)
1	(ABO), mRNA
NM 020445	Homo sapiens actin-related protein 3-beta (ARP3BETA), mRNA
NM_020435	Homo sapiens connexin46.6 (CX46.6), mRNA
NM_020426	Homo sapiens lysozyme homolog (LOC57151), mRNA
NM_020379	Homo sapiens 1,2-alpha-mannosidase IC (HMIC), mRNA
NM_020407	Homo sapiens Rh type B glycoprotein (RHBG), mRNA

	(DDVII) DOVI
NM_020406	Homo sapiens polycythemia rubra vera 1; cell surface receptor (PRV1), mRNA
NM 020377	Homo sapiens cysteinyl leukotriene CysLT2 receptor; cDNA PSEC0146 from
	clone PLACE1006979 (LOC57105), mRNA
NM_020355	Homo sapiens HRPAP20 short form (LOC57090), mRNA
NM 020350	Homo sapiens ATRAP protein (ATRAP), mRNA
NM 020380	Homo sapiens AF15q14 protein (AF15Q14), mRNA
NM_020368	Homo sapiens disrupter of silencing 10 (SAS10), mRNA
NM 020344	Homo sapiens solute carrier family 24 (sodium/potassium/calcium exchanger),
_	member 2 (SLC24A2), mRNA
NM_020396	Homo sapiens BCL2-like 10 (apoptosis facilitator) (BCL2L10), mRNA
NM_020384	Homo sapiens claudin 2 (CLDN2), mRNA
NM 007260	Homo sapiens lysophospholipase II (LYPLA2), mRNA
NM 000390	Homo sapiens choroideremia (Rab escort protein 1) (CHM), mRNA
NM 001994	Homo sapiens coagulation factor XIII, B polypeptide (F13B), mRNA
NM 000129	Homo sapiens coagulation factor XIII, A1 polypeptide (F13A1), mRNA
NM 000505	Homo sapiens coagulation factor XII (Hageman factor) (F12), mRNA
NM 000504	Homo sapiens coagulation factor X (F10), mRNA
NM_005509	Homo sapiens Dmx-like 1 (DMXL1), mRNA
NM 001300	Homo sapiens core promoter element binding protein (COPEB), mRNA
NM 012089	Homo sapiens ATP-binding cassette, sub-family B (MDR/TAP), member 10
_	(ABCB10), nuclear gene encoding mitochondrial protein, mRNA
NM 007188	Homo sapiens ATP-binding cassette, sub-family B (MDR/TAP), member 8
-	(ABCB8), nuclear gene encoding mitochondrial protein, mRNA
NM 005689	Homo sapiens ATP-binding cassette, sub-family B (MDR/TAP), member 6
_	(ABCB6), nuclear gene encoding mitochondrial protein, mRNA
NM 001216	Homo sapiens carbonic anhydrase IX (CA9), mRNA
NM_000717	Homo sapiens carbonic anhydrase IV (CA4), mRNA
NM_001218	Homo sapiens carbonic anhydrase XII (CA12), mRNA
NM_001217	Homo sapiens carbonic anhydrase XI (CA11), mRNA
NM_006384	Homo sapiens calcium and integrin binding protein (DNA-dependent protein
	kinase interacting protein) (SIP2-28), mRNA
NM_016734	Homo sapiens paired box gene 5 (B-cell lineage specific activator protein)
	(PAX5), mRNA
NM_000687	Homo sapiens S-adenosylhomocysteine hydrolase (AHCY), mRNA
NM_004482	Homo sapiens UDP-N-acetyl-alpha-D-galactosamine:polypeptide N-
	acetylgalactosaminyltransferase 3 (GalNAc-T3) (GALNT3), mRNA
NM_004481	Homo sapiens UDP-N-acetyl-alpha-D-galactosamine:polypeptide N-
	acetylgalactosaminyltransferase 2 (GalNAc-T2) (GALNT2), mRNA
NM_000512	Homo sapiens galactosamine (N-acetyl)-6-sulfate sulfatase (Morquio syndrome,
	mucopolysaccharidosis type IVA) (GALNS), mRNA
NM_000403	Homo sapiens galactose-4-epimerase, UDP- (GALE), mRNA
NM_020310	Homo sapiens MAX binding protein (MNT), mRNA
NM_006250	Homo sapiens proline-rich protein Haell subfamily 1 (PRH1), mRNA
NM_005164	Homo sapiens ATP-binding cassette, sub-family D (ALD), member 2 (ABCD2),
177.6.653555	mRNA
NM_020300	Homo sapiens microsomal glutathione S-transferase 1 (MGST1), mRNA
NM_000728	Homo sapiens calcitonin-related polypeptide, beta (CALCB), mRNA
NM_020127	Homo sapiens tuftelin 1 (TUFT1), mRNA
NM_020040	Homo sapiens tubulin, beta polypeptide 4, member Q (TUBB4Q), mRNA
NM_020126	Homo sapiens sphingosine kinase type 2 isoform (SPHK2), mRNA
NM_020203	Homo sapiens matrix, extracellular phosphoglycoprotein with ASARM motif
L	(bone) (MEPE), mRNA

NM_020231	Homo sapiens x 010 protein (MDS010), mRNA
NM_020132	Homo sapiens lysophosphatidic acid acyltransferase-gamma1 (LPAAT-
NM_020132	gamma1), mRNA
NB4 020246	Homo sapiens cation-chloride cotransporter-interacting protein (LOC56996),
NM_020246	mRNA
ND4 020242	Homo sapiens mitochondrial import receptor Tom22 (LOC56993), mRNA
NM_020243	Homo sapiens non-kinase Cdc42 effector protein SPEC2 (LOC56990), mRNA
NM_020240	Homo sapiens ancient conserved domain protein 4 (LOC56939), mRNA
NM_020184	Homo sapiens Carbonic anhydrase-related protein 10 (LOC56934), mRNA
NM_020178	Homo sapiens Carbonic annydrase-related protein 10 (EOC30934), mRVA
NM_020155	Homo sapiens chromosome 11 hypothetical protein ORF4 (LOC56834), mRNA
NM_020179	Homo sapiens FN5 protein (FN5), mRNA
NM_020187	Homo sapiens DC12 protein (DC12), mRNA
NM_020156	Homo sapiens corel UDP-galactose:N-acetylgalactosamine-alpha-R beta 1,3-
	galactosyltransferase (C1GALT1), mRNA
NM_000352	Homo sapiens ATP-binding cassette, sub-family C (CFTR/MRP), member 8 (ABCC8), mRNA
NM_000374	Homo sapiens uroporphyrinogen decarboxylase (UROD), mRNA
NM_002872	Homo sapiens ras-related C3 botulinum toxin substrate 2 (rho family, small GTP
_	binding protein Rac2) (RAC2), mRNA
NM 004152	Homo sapiens ornithine decarboxylase antizyme 1 (OAZ1), mRNA
NM 002527	Homo sapiens neurotrophin 3 (NTF3), mRNA
NM 002295	Homo sapiens laminin receptor 1 (67kD, ribosomal protein SA) (LAMR1),
	mRNA
NM 002293	Homo sapiens laminin, gamma 1 (formerly LAMB2) (LAMC1), mRNA
NM 002292	Homo sapiens laminin, beta 2 (laminin S) (LAMB2), mRNA
NM 002290	Homo sapiens laminin, alpha 4 (LAMA4), mRNA
NM 006192	Homo sapiens paired box gene 1 (PAX1), mRNA
NM 019896	Homo sapiens DNA polymerase epsilon p12 subunit (P12), mRNA
NM_000583	Homo sapiens group-specific component (vitamin D binding protein) (GC), mRNA
NM_019891	Homo sapiens endoplasmic reticulum oxidoreductin 1-Lbeta (ERO1-L(BETA)),
	mRNA
NM_006705	Homo sapiens growth arrest and DNA-damage-inducible, gamma (GADD45G), mRNA
NM_001924	Homo sapiens growth arrest and DNA-damage-inducible, alpha (GADD45A), mRNA
NM_019844	Homo sapiens solute carrier family 21 (organic anion transporter), member 8 (SLC21A8), mRNA
NM_019644	Homo sapiens testis-specific ankyrin motif containing protein (LOC56311), mRNA
NM_019842	Homo sapiens potassium voltage-gated channel, KQT-like subfamily, member 5 (KCNQ5), mRNA
NM_012281	Homo sapiens potassium voltage-gated channel, Shal-related subfamily, member 2 (KCND2), mRNA
NM_019857	Homo sapiens CTP synthase II (CTPS2), mRNA
NM_019839	Homo sapiens seven transmembrane receptor BLTR2; leukotriene B4 receptor
NR 005757	BLT2 (BLTR2), mRNA
NM_005757	Homo sapiens C3H-type zinc finger protein; similar to D. melanogaster muscleblind B protein (MBLL), mRNA
NM_004299	Homo sapiens ATP-binding cassette, sub-family B (MDR/TAP), member 7 (ABCB7), nuclear gene encoding mitochondrial protein, mRNA
NM 004683	Homo sapiens regucalcin (senescence marker protein-30) (RGN), mRNA

NM_019618	Homo sapiens interleukin-1 homolog 1 (IL-1H1), mRNA
NM 018950	Homo sapiens major histocompatibility complex, class I, F (HLA-F), mRNA
NM 019610	Homo sapiens hypothetical protein 669 (LOC56267), mRNA
NM 000523	Homo sapiens homeo box D13 (HOXD13), mRNA
NM 019607	Homo sapiens hypothetical protein FLJ11267 (FLJ11267), mRNA
NM 019604	Homo sapiens class-I MHC-restricted T cell associated molecule (CRTAM),
	mRNA
NM 012328	Homo sapiens microvascular endothelial differentiation gene 1 (MDG1), mRNA
NM 013303	Homo sapiens fetal hypothetical protein (HSU84971), mRNA
NM 013298	Homo sapiens hypothetical protein (HSU79252), mRNA
NM 013386	Homo sapiens hypothetical protein (DKFZp586G0123), mRNA
NM 013313	Homo sapiens hypothetical protein (AF060862), mRNA
NM 019116	Homo sapiens similar to ubiquitin binding protein (UBPH), mRNA
NM 018961	Homo sapiens ubiquitin associated and SH3 domain containing, A (UBASH3A),
14141_010301	mRNA
NM 018968	Homo sapiens syntrophin, gamma 2 (SNTG2), mRNA
NM 018967	Homo sapiens syntrophin, gamma 1 (SNTG1), mRNA
NM 018969	Homo sapiens super conserved receptor expressed in brain 3 (SREB3), mRNA
NM 018964	Homo sapiens solute carrier family 37 (glycerol-3-phosphate transporter),
14141_019304	member 1 (SLC37A1), mRNA
NIM OTODAS	Homo sapiens phosphodiesterase 7B (PDE7B), mRNA
NM_018945	Homo sapiens MAGE-like 2 (MAGEL2), mRNA
NM_019066	Homo sapiens NICE-1 protein (NICE-1), mRNA
NM_019060	Homo sapiens NICE-1 protein (NICE-1), INKNA Homo sapiens hypothetical protein (LOC55924), mRNA
NM_019099	Homo sapiens hypothetical protein (LOC53524), mixtx
NM_019003	Homo sapiens spindlin-like (LOC54466), mRNA
NM_018952	Homo sapiens homeo box B6 (HOXB6), mRNA
NM_018951	Homo sapiens homeo box A10 (HOXA10), mRNA
NM_018942	Homo sapiens homeo box (H6 family) 1 (HMX1), mRNA
NM_019109	Homo sapiens beta-1,4 mannosyltransferase (HMT-1), mRNA
NM_019052	Homo sapiens HCR (a-helix coiled-coil rod homologue) (HCR), mRNA
NM_018985	Homo sapiens hypothetical protein (HCGIV.9), mRNA
NM_019096	Homo sapiens GTP binding protein 2 (GTPBP2), mRNA
NM_018949	Homo sapiens G protein-coupled receptor 14 (GPR14), mRNA
NM_019048	Homo sapiens hypothetical protein (FLJ20752), mRNA
NM_019086	Homo sapiens hypothetical protein FLJ20674 (FLJ20674), mRNA
NM_019040	Homo sapiens hypothetical protein (FLJ20498), mRNA
NM_018988	Homo sapiens hypothetical protein (FLJ20330), mRNA
NM_019005	Homo sapiens hypothetical protein (FLJ20323), mRNA
NM_019027	Homo sapiens hypothetical protein (FLJ20273), mRNA
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NM 019000	Homo sapiens hypothetical protein (FLJ20152), mRNA
NM_019087	Homo sapiens hypothetical protein FLJ20051 (FLJ20051), mRNA
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NM_019021	Homo sapiens hypothetical protein (FLJ20010), mRNA
NM 019018	Homo sapiens hypothetical protein (FLJ11127), mRNA
NM 019084	Homo sapiens hypothetical protein FLJ10895 (FLJ10895), mRNA
NM 019070	Homo sapiens hypothetical protein (FLJ10432), mRNA
NM 019088	Homo sapiens hypothetical protein F23149_1 (F23149_1), mRNA
NM 019002	Homo sapiens ETAA16 protein (ETAA16), mRNA
NM 019114	Homo sapiens EHM2 gene (EHM2), mRNA
	Homo sapiens dolichyl-phosphate mannosyltransferase polypeptide 3 (DPM3),
NM_018973	mRNA
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27.5.010050	Homo sapiens DAZ associated protein 1 (DAZAP1), mRNA
NM_018959	Homo sapiens DAZ associated protein 1 (DAZAT 1), inicita
NM_019098	Homo sapiens cyclic nucleotide gated channel beta 3 (CNGB3), mRNA
NM_018958	Homo sapiens chromosome 15 open reading frame 2 (C15ORF2), mRNA
NM_000379	Homo sapiens xanthene dehydrogenase (XDH), mRNA
NM_000552	Homo sapiens von Willebrand factor (VWF), mRNA
NM_000362	Homo sapiens tissue inhibitor of metalloproteinase 3 (Sorsby fundus dystrophy,
	pseudoinflammatory) (TIMP3), mRNA
NM 003255	Homo sapiens tissue inhibitor of metalloproteinase 2 (TIMP2), mRNA
NM_003001	Homo sapiens succinate dehydrogenase complex, subunit C, integral membrane
-	protein, 15kD (SDHC), nuclear gene encoding mitochondrial protein, mRNA
NM_003000	Homo sapiens succinate dehydrogenase complex, subunit B, iron sulfur (Ip)
	(SDHB), nuclear gene encoding mitochondrial protein, mRNA
NM 006745	Homo sapiens sterol-C4-methyl oxidase-like (SC4MOL), mRNA
NM 006860	Homo sapiens putative GTP-binding protein similar to RAY/RAB1C (RAYL),
14147_000000	mRNA
NM 000531	Homo sapiens ornithine carbamoyltransferase (OTC), nuclear gene encoding
11111_000551	mitochondrial protein, mRNA
NM 000607	Homo sapiens orosomucoid 1 (ORM1), mRNA
NM 002538	Homo sapiens occludin (OCLN), mRNA
NM 002301	Homo sapiens lactate dehydrogenase C (LDHC), transcript variant 1, mRNA
NM 017448	Homo sapiens lactate dehydrogenase C (LDHC), transcript variant 2, mRNA
	Homo sapiens kallikrein B, plasma (Fletcher factor) 1 (KLKB1), mRNA
NM_000892	Homo sapiens inhibin, beta B (activin AB beta polypeptide) (INHBB), mRNA
NM_002193	
NM_002191	Homo sapiens inhibin, alpha (INHA), mRNA
NM_002015	Homo sapiens forkhead box O1A (rhabdomyosarcoma) (FOXO1A), mRNA
NM_004473	Homo sapiens forkhead box E1 (thyroid transcription factor 2) (FOXE1), mRNA
NM_000804	Homo sapiens folate receptor 3 (gamma) (FOLR3), mRNA
NM_000803	Homo sapiens folate receptor 2 (fetal) (FOLR2), mRNA
NM_004742	Homo sapiens BAI1-associated protein 1 (BAIAP1), mRNA
NM_004925	Homo sapiens aquaporin 3 (AQP3), mRNA
NM_007182	Homo sapiens Ras association (RalGDS/AF-6) domain family 1 (RASSF1),
	mRNA
NM_018941	Homo sapiens ceroid-lipofuscinosis, neuronal 8 (epilepsy, progressive with
	mental retardation) (CLN8), mRNA
NM_016936	Homo sapiens ubinuclein 1 (UBN1), mRNA
NM_012406	Homo sapiens PR domain containing 4 (PRDM4), mRNA
NM 018728	Homo sapiens myosin 5C (MYO5C), mRNA
NM_017540	Homo sapiens hypothetical protein DKFZp586H0623 (DKFZp586H0623),
_	mRNA
NM_018651	Homo sapiens zinc finger protein (ZFP), mRNA
NM_017503	Homo sapiens surfeit 2 (SURF2), mRNA
NM 018419	Homo sapiens SRY (sex determining region Y)-box 18 (SOX18), mRNA
NM_018427	Homo sapiens RNA polymerase I transcription factor RRN3 (RRN3), mRNA
NM_018545	Homo sapiens hypothetical protein PRO2955 (PRO2955), mRNA
NM 018525	Homo sapiens hypothetical protein PRO2369 (PRO2369), mRNA
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NM_018605	Homo sapiens hypothetical protein PRO1777 (PRO1777), mRNA
NM 018573	Homo sapiens hypothetical protein PRO1068 (PRO1068), mRNA
	Homo sapiens hypothetical protein PRO1051 (PRO1051), mRNA
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NM_018592	Homo sapiens hypothetical protein PRO0800 (PRO0800), mRNA
NM_018563	Homo sapiens hypothetical protein PRO0758 (PRO0758), mRNA

mo sapiens PR domain containing 5 (PRDM5), mRNA
mo sapiens myosin, heavy polypeptide 2, skeletal muscle, adult (MYH2),
NA
mo sapiens uncharacterized hematopoietic stem/progenitor cells protein
OSO26 (MDSO26), mRNA
mo sapiens lipopolysaccharide specific response-7 protein (LSR7), mRNA
mo sapiens HSVI binding protein (LOC55913), mRNA
mo sapiens 22kDa peroxisomal membrane protein-like (LOC55895), mRNA
mo sapiens neuronal specific transcription factor DAT1 (LOC55885), mRNA
mo sapiens neuronal specific danscription factor BFTT (Boosses), industription factor
mo sapiens CS box-containing wD protein (EOC53884), micevit mo sapiens AD-012 protein (LOC55833), mRNA
mo sapiens AD-012 protein (EOC33633), micros mo sapiens potassium inwardly-rectifying channel, subfamily J, member 16
CNI16), mRNA
mo sapiens hypothetical protein (IRO039700), mRNA
mo sapiens hypothetical protein IMPACT (IMPACT), mRNA
mo sapiens FEV protein (HSRNAFEV), mRNA
mo sapiens leptin receptor gene-related protein (HSOBRGRP), mRNA
mo sapiens metaphase chromosome protein 1 (HSMCR30), mRNA
mo sapiens p65 protein (HSAJ2425), mRNA
mo sapiens hypothetical protein HDCMC04P (HDCMC04P), mRNA
omo sapiens hypothetical protein HDCGC21P (HDCGC21P), mRNA
omo sapiens hepatocellular carcinoma-associated antigen 66 (HCA66), mRNA
omo sapiens putative methyltransferase (HASJ4442), mRNA
omo sapiens hypothetical protein FLJ20837 (FLJ20837), mRNA
omo sapiens hypothetical protein FLJ20758 (FLJ20758), mRNA
omo sapiens hypothetical protein FLJ20707 (FLJ20707), mRNA
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omo sapiens hypothetical protein FLJ20216 (FLJ20216), mRNA
omo sapiens hypothetical protein FLJ20097 (FLJ20097), mRNA
omo sapiens hypothetical protein FLJ20070 (FLJ20070), mRNA

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NM 018317	Homo sapiens hypothetical protein FLJ11082 (FLJ11082), mRNA
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NM 018263	Homo sapiens hypothetical protein FLJ10898 (FLJ10898), mRNA
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NM 018202	Homo sapiens hypothetical protein FLJ10747 (FLJ10747), mRNA
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NM 018063	Homo sapiens hypothetical protein FLJ10339 (FLJ10339), mRNA
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	Homo sapiens hypothetical protein FLJ10083 (FLJ10083), mRNA
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	Homo sapiens hypothetical protein FLJ10043 (FLJ10043), mRNA
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	Homo sapiens hypothetical protein FLJ10034 (FLJ10034), mRNA
NM 017973	Homo sapiens hypothetical protein DKFZp761D081 (DKFZp761D081), mRNA
NM 017610	Homo sapiens DKFZp564J157 protein (DKFZP564J157), mRNA
NM_018457	Homo sapiens DKFZp3643137 protein (DKFZp434K0920 (DKFZp434K0920),
NM_017590	mRNA
NM_017566	Homo sapiens hypothetical protein DKFZp434G0522 (DKFZp434G0522), mRNA
NM_017612	Homo sapiens hypothetical protein DKFZp434E2220 (DKFZp434E2220),
_	mRNA
NM_018641	Homo sapiens chondroitin 4-O-sulfotransferase 2 (C4S-2), mRNA
NM 018659	Homo sapiens cytokine-like protein C17 (C17), mRNA

NB/ 010656	Homo sapiens bladder cancer overexpressed protein (BLOV1), mRNA
NM_018656	Homo sapiens double-stranded RNA specific adenosine deaminase (ADAR3),
NM_018702	mRNA
ND4 014160	Homo sapiens HSPC070 protein (HSPC070), mRNA
NM_014160 NM_004288	Homo sapiens pleckstrin homology, Sec7 and coiled/coil domains, binding
NM_004288	protein (PSCDBP), mRNA
NM 004060	Homo sapiens cyclin G1 (CCNG1), mRNA
	Homo sapiens transcription factor binding to IGHM enhancer 3 (TFE3), mRNA
NM_006521 NM_007035	Homo sapiens keratocan (KERA), mRNA
	Homo sapiens tumor protein p53 (Li-Fraumeni syndrome) (TP53), mRNA
NM_000546	Homo sapiens secreted frizzled-related protein 5 (SFRP5), mRNA
NM_003015	Homo sapiens secreted frizzled-related protein 1 (SFRP1), mRNA
NM_003012	Homo sapiens ubiquitin specific protease 18 (USP18), mRNA
NM_017414	Homo sapiens ubiquitin associated protein (UBAP), mRNA
NM_016525	Homo sapiens toll-like receptor 9 (TLR9), mRNA
NM_017442	Homo sapiens polymerase (DNA directed), alpha (POLA), mRNA
NM_016937	Homo sapiens NADPH oxidase 4 (NOX4), mRNA
NM 016931	Homo sapiens myosin IIIA (MYO3A), mRNA
NM 017433	Homo sapiens junctional adhesion molecule (JAM), mRNA
NM_016946	Homo sapiens inositol(myo)-1(or 4)-monophosphatase 1 (IMPA1), mRNA
NM_005536 NM_017410	Homo sapiens homeo box C13 (HOXC13), mRNA
	Homo sapiens homeo box C10 (HOXC10), mRNA
NM_017409	Homo sapiens NAD(P) dependent steroid dehydrogenase-like; H105e3
NM_015922	(H105E3), mRNA
NM 004129	Homo sapiens guanylate cyclase 1, soluble, beta 2 (GUCY1B2), mRNA
NM 017423	Homo sapiens UDP-N-acetyl-alpha-D-galactosamine:polypeptide N-
10101_017423	acetylgalactosaminyltransferase 7 (GalNAc-T7) (GALNT7), mRNA
NM 016947	Homo sapiens G8 protein (G8), mRNA
NM_017434	Homo sapiens dual oxidase 1 (DUOX1), mRNA
NM 012143	Homo sapiens tuftelin-interacting protein (TIP39), mRNA
NM 017418	Homo sapiens deleted in esophageal cancer 1 (DEC1), mRNA
NM 016929	Homo sapiens chloride intracellular channel 5 (CLIC5), mRNA
NM 017413	Homo sapiens apelin; peptide ligand for APJ receptor (APELIN), mRNA
NM 000477	Homo sapiens albumin (ALB), mRNA
NM_007235	Homo sapiens exportin, tRNA (nuclear export receptor for tRNAs) (XPOT),
****	mRNA
NM_004585	Homo sapiens retinoic acid receptor responder (tazarotene induced) 3
1	(RARRES3), mRNA
NM 002134	Homo sapiens heme oxygenase (decycling) 2 (HMOX2), mRNA
NM 002100	Homo sapiens glycophorin B (includes Ss blood group) (GYPB), mRNA
NM 002099	Homo sapiens glycophorin A (includes MN blood group) (GYPA), mRNA
NM 005708	Homo sapiens glypican 6 (GPC6), mRNA
NM_013280	Homo sapiens fibronectin leucine rich transmembrane protein 1 (FLRT1),
	mRNA
NM_001304	Homo sapiens carboxypeptidase D (CPD), mRNA
NM_013410	Homo sapiens adenylate kinase 3 (AK3), nuclear gene encoding mitochondrial
	protein, mRNA
NM_002161	Homo sapiens isoleucine-tRNA synthetase (IARS), transcript variant short,
	mRNA
NM_013417	Homo sapiens isoleucine-tRNA synthetase (IARS), transcript variant long,
	mRNA
NM_015836	Homo sapiens tryptophanyl tRNA synthetase 2 (mitochondrial) (WARS2),

1' mite de de la mateira mPNA
nuclear gene encoding mitochondrial protein, mRNA
Homo sapiens methyl CpG binding protein 2 (Rett syndrome) (MECP2), mRNA
Homo sapiens methyl-CpG binding domain protein 3 (MBD3), mRNA
Homo sapiens LIM domain only 6 (LMO6), mRNA
Homo sapiens killer cell lectin-like receptor subfamily C, member 4 (KLRC4),
mRNA
Homo sapiens engrailed homolog 2 (EN2), mRNA
Homo sapiens engrailed homolog 1 (EN1), mRNA
Homo sapiens zinc finger protein 155 (pHZ-96) (ZNF155), mRNA
Homo sapiens zinc finger protein (ZFD25) (ZFD25), mRNA
Homo sapiens Wilms tumor associated protein (WIT-1), mRNA
Homo sapiens villin-like (VILL), mRNA
Homo sapiens variable charge protein on X with eight repeats (VCX-8r), mRNA
Homo sapiens variable charge protein on X with two repeats (VCX-2r), mRNA
Homo sapiens tubulin, gamma 2 (TUBG2), mRNA
Homo sapiens TU12B1-TY protein (TU12B1-TY), mRNA
Homo sapiens KRAB-zinc finger protein SZF1-1 (SZF1), mRNA
Homo sapiens solute carrier family 21 (organic anion transporter), member 11
(SLC21A11), mRNA
Homo sapiens putative secreted protein (SIG11), mRNA
Homo sapiens SH3 and PX domain-containing protein SH3PX1 (SH3PX1),
mRNA PAGENTAL PAGENTA
Homo sapiens serum/glucocorticoid regulated kinase 2 (SGK2), mRNA
Homo sapiens S2P protein (S2P), mRNA
Homo sapiens RU2S (RU2), mRNA
Homo sapiens Rh type C glycoprotein (RHCG), mRNA
Homo sapiens phosphatidylserine-specific phospholipase Alalpha (PS-PLA1),
mRNA
Homo sapiens ninjurin 2 (NINJ2), mRNA
Homo sapiens membrane interacting protein of RGS16 (MIR16), mRNA
Homo sapiens integral inner nuclear membrane protein (MAN1), mRNA
Homo sapiens melanoma antigen, family E, 1, cancer/testis specific (MAGEE1),
mRNA
Homo sapiens LW-1 (LW-1), mRNA
Homo sapiens seven transmembrane protein TM7SF3 (TM7SF3), mRNA
Homo sapiens ATPase, aminophospholipid transporter-like, Class I, type 8A,
member 2 (ATP8A2), mRNA
Homo sapiens synoretin (LOC51749), mRNA
Homo sapiens ghrelin precursor (LOC51738), mRNA
Homo sapiens Kruppel-like factor (LOC51713), mRNA
Homo sapiens cytochrome b5 reductase 1 (B5R.1) (LOC51706), mRNA
Homo sapiens nemo-like kinase (LOC51701), mRNA
Homo sapiens RhD type IIIa protein (LOC51698), mRNA
Homo sapiens alpha 1,2-mannosidase (LOC51697), mRNA
Homo sapiens hHDC for homolog of Drosophila headcase (LOC51696), mRNA
Homo sapiens U6 snRNA-associated Sm-like protein LSm7 (LOC51690),
mRNA
Homo sapiens prothymosin a14 (LOC51685), mRNA
Homo sapiens MAGUK protein p55T; Protein Associated with Lins 2
(LOC51678), mRNA
Homo sapiens HSPCO34 protein (LOC51668), mRNA
Homo sapiens NY-REN-18 antigen (LOC51667), mRNA

NM_016079	Homo sapiens CGI-149 protein (LOC51652), mRNA
NM_016062	Homo sapiens CGI-128 protein (LOC51647), mRNA
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NM 016016	Homo sapiens CGI-69 protein (LOC51629), mRNA
NM 016008	Homo sapiens CGI-60 protein (LOC51626), mRNA
NM 015995	Homo sapiens Kruppel-like factor 13 (KLF13), mRNA
NM 015980	Homo sapiens HMP19 protein (LOC51617), mRNA
NM 015958	Homo sapiens CGI-30 protein (LOC51611), mRNA
NM 015941	Homo sapiens CGI-11 protein (LOC51606), mRNA
NM 015937	Homo sapiens CGI-06 protein (LOC51604), mRNA
NM 015929	Homo sapiens lipoyltransferase (LOC51601), mRNA
NM 015921	Homo sapiens divalent cation tolerant protein CUTA (LOC51596), mRNA
NM 015908	Homo sapiens arsenate resistance protein ARS2 (ARS2), mRNA
NM 015875	Homo sapiens unnamed HERV-H protein (LOC51581), mRNA
NM 015874	Homo sapiens H-2K binding factor-2 (LOC51580), mRNA
NM 016283	Homo sapiens adrenal gland protein AD-004 (LOC51578), mRNA
NM 016644	Homo sapiens mesenchymal stem cell protein DSC54 (LOC51334), mRNA
NM 016643	Homo sapiens mesenchymal stem cell protein DSC43 (LOC51333), mRNA
NM 016642	Homo sapiens beta V spectrin (BSPECV), mRNA
NM 016638	Homo sapiens SRp25 nuclear protein (LOC51329), mRNA
NM 016637	Homo sapiens neaml (LOC51328), mRNA
NM 016633	Homo sapiens EDRF protein (LOC51327), mRNA
NM 016625	Homo sapiens hypothetical protein (LOC51319), mRNA
NM 016622	Homo sapiens hypothetical protein (LOC51318), mRNA
NM 016621	Homo sapiens hypothetical protein (LOC51317), mRNA
NM 016609	Homo sapiens hBOIT for potent brain type organic ion transporter (LOC51310),
1411_01000>	mRNA
NM 016606	Homo sapiens SGC32445 protein (LOC51308), mRNA
NM 016591	Homo sapiens core 2 beta-1,6-N-acetylglucosaminyltransferase 3 (LOC51301),
	mRNA
NM 016585	Homo sapiens testicular haploid expressed gene (THEG), mRNA
NM_016573	Homo sapiens Gem-interacting protein (LOC51291), mRNA
NM_016568	Homo sapiens G-protein coupled receptor SALPR; somatostatin and angiotensin-
	like peptide receptor (LOC51289), mRNA
NM 016566	Homo sapiens pparl (LOC51288), mRNA
NM_016563	Homo sapiens Ris (LOC51285), mRNA
NM_016548	Homo sapiens golgi membrane protein GP73 (LOC51280), mRNA
NM 016499	Homo sapiens hypothetical protein (LOC51259), mRNA
NM_016490	Homo sapiens hypothetical protein (LOC51252), mRNA
NM 016466	Homo sapiens hypothetical protein (LOC51239), mRNA
NM 016459	Homo sapiens hypothetical protein (LOC51237), mRNA
NM 016449	Homo sapiens hypothetical protein (LOC51233), mRNA
NM 016440	Homo sapiens VRK3 for vaccinia related kinase 3 (LOC51231), mRNA
NM 016427	Homo sapiens transcription elongation factor (SIII) elongin A2 (TCEB3L),
	mRNA
NM 016423	Homo sapiens zinc finger protein 219 (ZNF219), mRNA
NM_016361	Homo sapiens LPAP for lysophosphatidic acid phosphatase (LOC51205),
<u>-</u>	mRNA
NM_016353	Homo sapiens r c (LOC51201), mRNA
NM_016349	Homo sapiens susceptibility protein NSG-x (LOC51198), mRNA

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NM_016341	Homo sapiens pancreas-enriched phospholipase C (LOC51196), mRNA
NM_016323	Homo sapiens cyclin-E binding protein 1 (LOC51191), mRNA
NM_016317	Homo sapiens neutral sphingomyelinase (LOC51190), mRNA
NM_016286	Homo sapiens carbonyl reductase (LOC51181), mRNA
NM_016269	Homo sapiens lymphoid enhancer binding factor-1 (LOC51176), mRNA
NM_016245	Homo sapiens retinal short-chain dehydrogenase/reductase retSDR2
	(LOC51170), mRNA
NM 016241	Homo sapiens endomucin-1 (LOC51169), mRNA
NM 016230	Homo sapiens flavohemoprotein b5+b5R (LOC51167), mRNA
NM 016221	Homo sapiens dynactin p62 subunit (LOC51164), mRNA
NM 016215	Homo sapiens NEU1 protein (LOC51162), mRNA
NM 016210	Homo sapiens g20 protein (LOC51161), mRNA
NM 016161	Homo sapiens alpha-1,4-N-acetylglucosaminyltransferase (LOC51146), mRNA
NM 016123	Homo sapiens putative protein kinase NY-REN-64 antigen (LOC51135), mRNA
NM 016120	Homo sapiens putative ring zinc finger protein NY-REN-43 antigen
	(LOC51132), mRNA
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NM 016011	Homo sapiens CGI-63 protein (LOC51102), mRNA
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NM 015999	Homo sapiens CGI-45 protein (LOC51094), mRNA
NM 015982	Homo sapiens germ cell specific Y-box binding protein (LOC51087), mRNA
NM 015963	Homo sapiens CGI-36 protein (LOC51078), mRNA
NM 015959	Homo sapiens CGI-31 protein (LOC51075), mRNA
NM 015950	Homo sapiens CGI-22 protein (LOC51069), mRNA
NM 015938	Homo sapiens CGI-07 protein (LOC51068), mRNA
NM 015916	Homo sapiens hypothetical protein (LOC51063), mRNA
NM 015914	Homo sapiens hypothetical protein (LOC51061), mRNA
NM 015910	Homo sapiens hypothetical protein (LOC51057), mRNA
NM 015901	Homo sapiens unknown (LOC51055), mRNA
NM 015893	Homo sapiens preproprolactin-releasing peptide (LOC51052), mRNA
NM_015887	Homo sapiens putative peroxisome microbody protein 175.1 (LOC51051),
	mRNA
NM 015880	Homo sapiens RIG-like 14-1 (LOC51047), mRNA
NM 015877	Homo sapiens Kruppel-associated box protein (LOC51045), mRNA
NM 015863	Homo sapiens surfactant protein B (LOC51041), mRNA
NM 015854	Homo sapiens retinoic acid receptor-beta associated open reading frame
	(LOC51036), mRNA
NM 015849	Homo sapiens pancreatic elastase IIB (LOC51032), mRNA
NM 016075	Homo sapiens CGI-145 protein (LOC51028), mRNA
NM 016074	Homo sapiens CGI-143 protein (LOC51027), mRNA
NM 016063	Homo sapiens CGI-130 protein (LOC51020), mRNA
NM 016048	Homo sapiens CGI-111 protein (LOC51015), mRNA
NM 016044	Homo sapiens CGI-105 protein (LOC51011), mRNA
NM 015947	Homo sapiens CGI-18 protein (LOC51008), mRNA
NM 016058	Homo sapiens CGI-121 protein (LOC51002), mRNA
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	COLIO (COCCIONO)
NM_015948	Homo sapiens CGI-19 protein (LOC51000), mRNA
NM_016040	Homo sapiens CGI-100 protein (LOC50999), mRNA
NM_016571	Homo sapiens lengsin (LGS), mRNA
NM_015868	Homo sapiens NK-receptor (KIR-023GB), mRNA
NM_016281	Homo sapiens STE20-like kinase (JIK), mRNA
NM_016358	Homo sapiens iroquois homeobox protein 4 (IRX4), mRNA
NM_016291	Homo sapiens mammalian inositol hexakisphosphate kinase 2 (IP6K2), mRNA
NM_015848	Homo sapiens cytokeratin 2 (HUMCYT2A), mRNA
NM_016506	Homo sapiens hypothetical protein (HSPC252), mRNA
NM_016498	Homo sapiens hypothetical protein (HSPC242), mRNA
NM_016460	Homo sapiens hypothetical protein (HSPC192), mRNA
NM_016390	Homo sapiens hypothetical protein (HSPC109), mRNA
NM_016091	Homo sapiens HSPC025 (HSPC025), mRNA
NM_016522	Homo sapiens neurotrimin (HNT), mRNA
NM_016258	Homo sapiens high-glucose-regulated protein 8 (HGRG8), mRNA
NM_016173	Homo sapiens HEMK homolog 7kb (HEMK), mRNA
NM_016516	Homo sapiens tumor antigen SLP-8p (HCC8), mRNA
NM_016540	Homo sapiens G protein-coupled receptor 72 (GPR72), mRNA
NM_012196	Homo sapiens G antigen 8 (GAGE8), mRNA
NM_015898	Homo sapiens HIV-1 inducer of short transcripts binding protein (FBI1), mRNA
NM_016357	Homo sapiens epithelial protein lost in neoplasm beta (EPLIN), mRNA
NM_016218	Homo sapiens polymerase (DNA-directed) kappa (POLK), mRNA
NM_016240_	Homo sapiens CSR1 protein (CSR1), mRNA
NM_016073	Homo sapiens CGI-142 (CGI-142), mRNA
NM_016315	Homo sapiens CED-6 protein (CED-6), mRNA
NM_016620	Homo sapiens hypothetical protein (BM-005), mRNA
NM_015896	Homo sapiens BLu protein (BLu), mRNA
NM_016426	Homo sapiens G-2 and S-phase expressed 1 (GTSE1), mRNA
NM_015928	Homo sapiens androgen-induced prostate proliferative shutoff associated protein (AS3), mRNA
NM 016238	Homo sapiens anaphase-promoting complex subunit 7 (APC7), mRNA
NM 016376	Homo sapiens ANKHZN protein (ANKHZN), mRNA
NM 016282	Homo sapiens adenylate kinase 3 alpha like (AKL3L), mRNA
NM 016453	Homo sapiens SH3 protein (AF3P21), mRNA
NM 016614	Homo sapiens TRAF and TNF receptor-associated protein (AD022), mRNA
NM_015365	Homo sapiens Alport syndrome, mental retardation, midface hypoplasia and
	elliptocytosis chromosomal region, gene 1 (AMMECR1), mRNA
NM_007126	Homo sapiens valosin-containing protein (VCP), mRNA
NM_001059	Homo sapiens tachykinin receptor 3 (TACR3), mRNA
NM_005963	Homo sapiens myosin, heavy polypeptide 1, skeletal muscle, adult (MYH1),
	mRNA PNA
NM_005561	Homo sapiens lysosomal-associated membrane protein 1 (LAMP1), mRNA
NM_006407	Homo sapiens vitamin A responsive; cytoskeleton related (JWA), mRNA
NM_000854	Homo sapiens glutathione S-transferase theta 2 (GSTT2), mRNA
NM_002046	Homo sapiens glyceraldehyde-3-phosphate dehydrogenase (GAPD), mRNA
NM_001953	Homo sapiens endothelial cell growth factor 1 (platelet-derived) (ECGF1), mRNA
NM 000927	Homo sapiens ATP-binding cassette, sub-family B (MDR/TAP), member 1
	(ABCB1), mRNA
NM_015686	Homo sapiens TED protein (TED), mRNA
NM 014070	
TATAT OT-1010	Homo sapiens STG protein (STG), mRNA Homo sapiens SPR1 protein (SPR1), mRNA

ND 6 014060	TI CEEVI mestein (CEEVI) mDMA
NM_014068	Homo sapiens SEEK1 protein (SEEK1), mRNA
NM_014051	Homo sapiens PTD011 protein (PTD011), mRNA
NM_014109	Homo sapiens PRO2000 protein (PRO2000), mRNA
NM_014107	Homo sapiens PRO1992 protein (PRO1992), mRNA
NM_014095	Homo sapiens PRO1600 protein (PRO1600), mRNA
NM_014084	Homo sapiens PRO0806 protein (PRO0806), mRNA
NM_014130	Homo sapiens PRO0483 protein (PRO0483), mRNA
NM_014082	Homo sapiens PRO0397 protein (PRO0397), mRNA
NM_014125	Homo sapiens PRO0327 protein (PRO0327), mRNA
NM_014081	Homo sapiens PRO0297 protein (PRO0297), mRNA
NM_014037	Homo sapiens NTT5 protein (NTT5), mRNA
NM_015367	Homo sapiens MIL1 protein (MIL1), nuclear gene encoding mitochondrial
22.5.6.66	protein, mRNA
NM_014060	Homo sapiens MCT-1 protein (MCT-1), mRNA
NM_014892	Homo sapiens KIAA1116 protein (KIAA1116), mRNA
NM_014968	Homo sapiens KIAA1104 protein (KIAA1104), mRNA
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NM_014799	Homo sapiens hephaestin (HEPH), mRNA
NM_014678	Homo sapiens KIAA0685 gene product (KIAA0685), mRNA
NM_014011	Homo sapiens KIAA0671 gene product (KIAA0671), mRNA
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NM_014767	Homo sapiens KIAA0275 gene product (KIAA0275), mRNA
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NM_015153	Homo sapiens KIAA0244 protein (KIAA0244), mRNA
NM_014747	Homo sapiens KIAA0237 gene product (KIAA0237), mRNA
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NM 014640	Homo sapiens KIAA0173 gene product (KIAA0173), mRNA
NM 014666	Homo sapiens KIAA0171 gene product (KIAA0171), mRNA
NM 014641	Homo sapiens KIAA0170 gene product (KIAA0170), mRNA
NM_014737	Homo sapiens Ras association (RalGDS/AF-6) domain family 2 (RASSF2), mRNA
NM 014770	Homo sapiens KIAA0167 gene product (KIAA0167), mRNA
NM 014739	Homo saniens KIAA0164 gene product (KIAA0164), mRNA
NM_014865	Homo sapiens chromosome condensation-related SMC-associated protein 1
11112_01 1000	(KIAA0159), mRNA
NM 014748	Homo sapiens KIAA0064 gene product (KIAA0064), mRNA
NM 014876	Homo sapiens KIAA0063 gene product (KIAA0063), mRNA
NM 014764	Homo sapiens DAZ associated protein 2 (DAZAP2), mRNA
NM 014875	Homo sapiens KIAA0042 gene product (KIAA0042), mRNA
NM 014642	Homo sapiens KIAA0036 gene product (KIAA0036), mRNA
NM 015340	Homo sapiens leucyl-tRNA synthetase, mitochondrial (KIAA0028), mRNA
NM 014634	Homo sapiens KIAA0015 gene product (KIAA0015), mRNA
NM_014783	Homo sapiens KIAA0013 gene product (KIAA0013), mRNA
NM 014008	Homo sapiens JM1 protein (JM1), mRNA
NM 014066	Homo sapiens HT002 protein; hypertension-related calcium-regulated gene
14141_014000	(HT002), mRNA
NM_014154	Homo sapiens HSPC056 protein (HSPC056), mRNA
NM 014153	Homo sapiens HSPC055 protein (HSPC055), mRNA
NM 014150	Homo sapiens HSPC052 protein (HSPC052), mRNA
NM 014149	Homo sapiens HSPC049 protein (HSPC049), mRNA
NM 014029	Homo sapiens HSPC022 protein (HSPC022), mRNA
NM 014027	Homo sapiens HSPC018 protein (HSPC018), mRNA
NM 014019	Homo sapiens HSPC009 protein (HSPC009), mRNA
NM 015372	Homo sapiens hypothetical protein (HSN44A4A), mRNA
NM 015343	Homo sapiens hypothetical protein (HSA011916), mRNA
NM_014063	Homo sapiens src homology 3 domain-containing protein HIP-55 (HIP-55), mRNA
NM 014052	Homo sapiens GW128 protein (GW128), mRNA
NM 014888	Homo sapiens predicted osteoblast protein (GS3786), mRNA
NM 014030	Homo sapiens G protein-coupled receptor kinase-interactor 1 (GIT1), mRNA
NM 014077	Homo sapiens DKFZP586O0120 protein (DKFZP586O0120), mRNA
NM 015425	Homo sapiens DKFZP586M0122 protein (DKFZP586M0122), mRNA
NM 015456	Homo sapiens DKFZP586B0519 protein (DKFZP586B0519), mRNA
NM 015393	Homo sapiens DKFZP564O0823 protein (DKFZP564O0823), mRNA
NM 015421	Homo sapiens DKFZP564K2062 protein (DKFZP564K2062), mRNA
NM 015415	Homo sapiens DKFZP564B167 protein (DKFZP564B167), mRNA
NM 015527	Homo sapiens DKFZP434P1750 protein (DKFZP434P1750), mRNA
NM 015458	Homo sapiens DKFZP434K171 protein (DKFZP434K171), mRNA
NM 015599	Homo sapiens N-acetylglucosamine-phosphate mutase (AGM1), mRNA
NM 015434	Homo sapiens DKFZP434B168 protein (DKFZP434B168), mRNA
NM 015434 NM 015699	Homo sapiens hypothetical protein (DJ159A19.3), mRNA
NM_015697	Homo sapiens hypothetical protein (CL640), mRNA
	Homo sapiens hypothetical protein (CL25022), mRNA
NM_015702	1 Monto saprens hypometical protein (Consolut), materi

NM_015703	
	Homo sapiens CGI-96 protein (CGI-96), mRNA
NM_015380	Homo sapiens CGI-51 protein (CGI-51), mRNA
NM_014143	Homo sapiens B7-H1 protein (B7-H1), mRNA
NM_014062	Homo sapiens ART-4 protein (ART-4), mRNA
NM_014596	Homo sapiens zinc ribbon domain containing, 1 (ZNRD1), mRNA
NM_014519	Homo sapiens zinc finger protein 232 (ZNF232), mRNA
NM_014437	Homo sapiens zinc/iron regulated transporter-like (ZIRTL), mRNA
NM_015363	Homo sapiens zinc finger, imprinted 2 (ZIM2), mRNA
NM_014232	Homo sapiens vesicle-associated membrane protein 2 (synaptobrevin 2) (VAMP2), mRNA
NM_014233	Homo sapiens upstream binding transcription factor, RNA polymerase I (UBTF), mRNA
NM 014235	Homo sapiens ubiquitin-like 4 (UBL4), mRNA
NM 014383	Homo sapiens testis zinc finger protein (TZFP), mRNA
NM 014547	Homo sapiens tropomodulin 3 (ubiquitous) (TMOD3), mRNA
NM 014548	Homo sapiens tropomodulin 2 (neuronal) (TMOD2), mRNA
NM 014464	Homo sapiens tubulointerstitial nephritis antigen (TIN-AG), mRNA
NM 014258	Homo sapiens synaptonemal complex protein 2 (SYCP2), mRNA
NM 014370	Homo sapiens serine/threonine kinase 23 (STK23), mRNA
NM 014264	Homo sapiens serine/threonine kinase 18 (STK18), mRNA
NM 014467	Homo sapiens sushi-repeat protein (SRPUL), mRNA
NM 014230	Homo sapiens signal recognition particle 68kD (SRP68), mRNA
NM 014320	Homo sapiens putative heme-binding protein (SOUL), mRNA
NM 014426	Homo sapiens sorting nexin 5 (SNX5), mRNA
NM 014311	Homo sapiens single-strand selective monofunctional uracil DNA glycosylase
11111_014511	(SMUG1), mRNA
NM_014270	Homo sapiens solute carrier family 7 (cationic amino acid transporter, y+ system), member 9 (SLC7A9), mRNA
NM 014252	Homo sapiens solute carrier family 25 (mitochondrial carrier; ornithine
	transporter) member 15 (SLC25A15), nuclear gene encoding mitochondrial protein, mRNA
NM_014251	Homo sapiens solute carrier family 25, member 13 (citrin) (SLC25A13), mRNA
NM 014442	Homo sapiens sialic acid binding Ig-like lectin 8 (SIGLEC8), mRNA
NM 014521	Homo sapiens SH3-domain binding protein 4 (SH3BP4), mRNA
NM 014554	Homo sapiens sentrin/SUMO-specific protease (SENP1), mRNA
NM 014563	Homo sapiens spondyloepiphyseal dysplasia, late (SEDL), mRNA
NM 014191	Homo sapiens sodium channel, voltage gated, type VIII, alpha polypeptide
- 12.7207.12.71	(SCN8A), mRNA
NM_014139	Homo sapiens sodium channel, voltage-gated, type XII, alpha polypeptide (SCN12A), mRNA
NB4 014262	
NM_014363	Homo sapiens spastic ataxia of Charlevoix-Saguenay (sacsin) (SACS), mRNA
NM_014285	Homo sapiens homolog of Yeast RRP4 (ribosomal RNA processing 4), 3'-5'-exoribonuclease (RRP4), mRNA
NM_014496	Homo sapiens ribosomal protein S6 kinase, 90kD, polypeptide 6 (RPS6KA6), mRNA
NM 014245	Homo sapiens ring finger protein 7 (RNF7), mRNA
NM 014372	Homo sapiens ring finger protein 11 (RNF11), mRNA
NM 014314	Homo sapiens RNA helicase (RIG-I), mRNA
NM 014488	Homo sapiens RAB30, member RAS oncogene family (RAB30), mRNA
NM 014470 NM 014248 NM 014226	Homo sapiens GTP-binding protein (RHO6), mRNA Homo sapiens ring-box 1 (RBX1), mRNA Homo sapiens renal tumor antigen (RAGE), mRNA

NM 014353	Homo sapiens RAB26, member RAS oncogene family (RAB26), mRNA
NM 014410	Homo sapiens clusterin-like 1 (retinal) (CLUL1), mRNA
NM_015725	Homo sapiens photoreceptor outer segment all-trans retinol dehydrogenase
14141_013723	(PRRDH), mRNA
NM 005973	Homo sapiens papillary renal cell carcinoma (translocation-associated) (PRCC),
1111_003373	mRNA
NM_014337	Homo sapiens peptidylprolyl isomerase (cyclophilin)-like 2 (PPIL2), mRNA
NM_014348	Homo sapiens similar to rat integral membrane glycoprotein POM121
	(POM121L1), mRNA
NM_015720	Homo sapiens endoglycan (PODLX2), mRNA
NM_014386	Homo sapiens polycystic kidney disease 2-like 2 (PKD2L2), mRNA
NM_014390	Homo sapiens EBNA-2 co-activator (100kD) (p100), mRNA
NM_014321	Homo sapiens origin recognition complex, subunit 6 (yeast homolog)-like
	(ORC6L), mRNA
NM_014566	Homo sapiens olfactory receptor, family 1, subfamily D, member 5 (OR1D5), mRNA
NM 014565	Homo sapiens olfactory receptor, family 1, subfamily A, member 1 (OR1A1),
	mRNA
NM_014352	Homo sapiens POU transcription factor (OCT11), mRNA
NM_014581	Homo sapiens odorant-binding protein 2B (OBP2B), mRNA
NM_014582	Homo sapiens odorant-binding protein 2A (OBP2A), mRNA
NM_014142	Homo sapiens nudix (nucleoside diphosphate linked moiety X)-type motif 5
	(NUDT5), mRNA
NM_014502	Homo sapiens nuclear matrix protein NMP200 related to splicing factor PRP19
	(NMP200), mRNA
NM_014328	Homo sapiens nesca protein (NESCA), mRNA
NM_014222	Homo sapiens NADH dehydrogenase (ubiquinone) 1 alpha subcomplex, 8
ND 6 015679	(19kD, PGIV) (NDUFA8), mRNA
NM_015678	Homo sapiens neurobeachin (NBEA), mRNA Homo sapiens contactin 6 (CNTN6), mRNA
NM_014461 NM_014520	Homo sapiens MYB binding protein (P160) 1a (MYBBP1A), mRNA
NM 014221	Homo sapiens mature T-cell proliferation 1 (MTCP1), mRNA
NM 005927	Homo sapiens microfibrillar-associated protein 3 (MFAP3), mRNA
NM 014623	Homo sapiens male-enhanced antigen (MEA), mRNA
NM 014462	Homo sapiens Lsm1 protein (LSM1), mRNA
NM_014622	Homo sapiens loss of heterozygosity, 11, chromosomal region 2, gene A
11111_017022	(LOH11CR2A), mRNA
NM_014240	Homo sapiens LIM domains containing 1 (LIMD1), mRNA
NM_014564	Homo sapiens LIM homeobox protein 3 (LHX3), mRNA
NM_014553	Homo sapiens LBP protein (LBP-9), mRNA
NM_014387	Homo sapiens linker for activation of T cells (LAT), mRNA
NM_014379	Homo sapiens neuronal potassium channel alpha subunit (KV8.1), mRNA
NM_014514	Homo sapiens killer cell immunoglobulin-like receptor, three domains, short
	cytoplasmic tail, 1 (KIR3DS1), mRNA
NM_014513	Homo sapiens killer cell immunoglobulin-like receptor, two domains, short
	cytoplasmic tail, 5 (KIR2DS5), mRNA
NM_014512	Homo sapiens killer cell immunoglobulin-like receptor, two domains, short
27 6 01 4515	cytoplasmic tail, 1 (KIR2DS1), mRNA
NM_014511	Homo sapiens killer cell immunoglobulin-like receptor, two domains, long
ND 6 01 6010	cytoplasmic tail, 3 (KIR2DL3), mRNA
NM_014219	Homo sapiens killer cell immunoglobulin-like receptor, two domains, long
L	cytoplasmic tail, 2 (KIR2DL2), mRNA

ovtoplasmic tail, i (KIRZDL1), mRNA NM_014765 Homo sapiens translocase of outer mitochondrial membrane 20 (yeast) homolog (KIAA0016), mRNA NM_014406 NM_014406 NM_014407 Momo sapiens potassium large conductance calcium-activated channel, subfamily M, beta member 3-like (KCNMB3), mRNA NM_014407 NM_014407 NM_014407 NM_014408 NM_014416 Homo sapiens potassium large conductance calcium-activated channel, subfamily M, beta member 3-like (KCNMB3), mRNA NM_014216 Homo sapiens inositol 1,3,4-triphosphate 5/6 kinase (ITPK1), mRNA NM_014214 Homo sapiens inositol (INVS), mRNA NM_014214 Homo sapiens iniositol (INVS), mRNA NM_014211 Homo sapiens iniositol (INVS), mRNA NM_014231 Homo sapiens interleukin 17 receptor (IL17R), mRNA NM_014333 Homo sapiens interleukin 17B (IL17B), mRNA NM_014334 NM_014434 Homo sapiens interleukin 17B (IL17B), mRNA NM_014434 Homo sapiens phast shock 27kD protein B (HSU47926), mRNA NM_014434 Homo sapiens brant shock 27kD protein B (HSU47926), mRNA NM_01433 NM_01433 Homo sapiens heat shock 27kD protein family, member 7 (cardiovascular) (HSPB7), mRNA NM_01433 NM_01434 Homo sapiens bypothetical protein (HS347E2A), mRNA NM_01434 NM_01434 Homo sapiens bypothetical protein (HS347E2A), mRNA NM_01435 Homo sapiens bypothetical protein (HS347E2A), mRNA NM_01435 Homo sapiens bypothetical protein (HS74TE2A), mRNA NM_01435 Homo sapiens bypothetical protein (HS74TE2A), mRNA NM_01425 Homo sapiens brook on the transferase (HSA9761), mRNA NM_01425 Homo sapiens brook on the transferase (HSA9761), mRNA NM_01426 Homo sapiens brook on the transferase (HSA9761), mRNA NM_01427 Homo sapiens brook on the transferase (HSA9761), mRNA NM_01428 Homo sapiens brook on the transferase (CD2091), mRNA NM_01429 Homo sapiens brook on the transferase (RSA9761), mRNA NM_01436 Homo sapiens brook o	NM 014218	Homo sapiens killer cell immunoglobulin-like receptor, two domains, long
(KIAA0016), mRNA NM_014406 NM_014407 Homo sapiens potassium large conductance calcium-activated channel, subfamily M, beta member 3-like (KCNMB3L), mRNA NM_014216 Homo sapiens potassium large conductance calcium-activated channel, subfamily M beta member 3 (KCNMB3L), mRNA NM_014216 NM_014216 NM_014221 Homo sapiens inversin (INVS), mRNA NM_014211 Homo sapiens sinversin (INVS), mRNA NM_014211 Homo sapiens interleukin 1 receptor (IL.17R), mRNA NM_014211 Homo sapiens interleukin 17 receptor (IL.17R), mRNA NM_01433 Homo sapiens interleukin 17 receptor (IL.17R), mRNA NM_014433 Homo sapiens interleukin 17 receptor (IL.17R), mRNA NM_014434 Homo sapiens interleukin 17 receptor (IL.17R), mRNA NM_014434 Homo sapiens interleukin 17 receptor (IL.17R), mRNA NM_01433 Homo sapiens interleukin 17 receptor (IL.17R), mRNA NM_01433 Homo sapiens sinterleukin 17 receptor (IL.17R), mRNA NM_01434 Homo sapiens beat shock 27kD protein family, member 4 (IGSF4), mRNA NM_014462 Homo sapiens beat shock 27kD protein family, member 7 (cardiovascular) (HSPB7), mRNA NM_015370 Homo sapiens bypothetical protein (HS747E2A), mRNA NM_015371 Homo sapiens hypothetical protein (HS747E2A), mRNA NM_014345 Homo sapiens hypothetical protein (HS747E2A), mRNA NM_01435 Homo sapiens shore obox (11 (HOXC11), mRNA NM_01425 Homo sapiens shore obox (24 (HOXC4), mRNA NM_014213 Homo sapiens home obox (24 (HOXC4), mRNA NM_014210 Homo sapiens home obox (21 (HOXC11), mRNA NM_01435 Homo sapiens home obox (21 (HOXC11), mRNA NM_01450 Homo sapiens home obox (21 (HOXC11), mRNA NM_01450 Homo sapiens home obox (21 (HOXC11), mRNA NM_01451 Homo sapiens home obox (21 (HOXC11), mRNA NM_01451 Homo sapiens harrylenhancer-of-split related with YRPW motif-like (HEYL), mRNA NM_01457 Homo sapiens harrylenhancer-of-split related with YRPW motif-like (HEYL), mRNA NM_01457 Homo sapiens glutamate receptor, ionotropic, kainate 4 (GRIK4), mRNA NM_01457 Homo sapiens glutamate receptor, ionotropic, kainate 4 (GRIK4), mRNA NM_01457	1414_014210	
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NM 014213 Homo sapiens CD209 antigen-like (CD209L), mRNA	NM_014255	Homo sapiens transmembrane protein 4 (TMEM4), mRNA
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	NM_014364	Homo sapiens glyceraldehyde-3-phosphate dehydrogenase, testis-specific
	NM_015714	
		Homo sapiens FGF receptor activating protein 1 (FRAG1), mRNA

NM_014585	Homo sapiens solute carrier family 11 (proton-coupled divalent metal ion
27.6.44.44	transporters), member 3 (SLC11A3), mRNA Homo sapiens putative secreted ligand homologous to fjx1 (FJX1), mRNA
NM_014344	Homo sapiens Interleukin-1 Superfamily z (FIL1(ZETA)), mRNA
NM_014439	Homo sapiens Interleukin-1 Superfamily 2 (FIL1(ZETA)), mRNA Homo sapiens Interleukin-1 Superfamily 1 (FIL1(EPSILON)), mRNA
NM_014440	Homo sapiens Interleukin-1 Superfamily ((FIL1), mRNA
NM_014438	Homo sapiens interieurin-1 Superianiny e (FICI), interior
NM_014210	Homo sapiens ecotropic viral integration site 2A (EVI2A), mRNA
NM_014355	Homo sapiens enolase alpha, lung-specific (ENO1B), mRNA
NM_014600	Homo sapiens EH-domain containing 3 (EHD3), mRNA
NM_014601	Homo sapiens EH-domain containing 2 (EHD2), mRNA
NM_014503	Homo sapiens down-regulated in metastasis (DRIM), mRNA
NM_014549	Homo sapiens DKFZp434P211 protein (DKFZP434P211), mRNA
NM_014388	Homo sapiens novel putative protein similar to YIL091C yeast hypothetical 84
	kD protein from SGA1-KTR7 (DJ434O14.5), mRNA
NM_014618	Homo sapiens deleted in bladder cancer chromosome region candidate 1
	(DBCCR1), mRNA
NM_014392	Homo sapiens neuron-specific protein (D4S234E), mRNA
NM_004389	Homo sapiens catenin (cadherin-associated protein), alpha 2 (CTNNA2), mRNA
NM_014343	Homo sapiens claudin 15 (CLDN15), mRNA
NM_014887	Homo sapiens hypothetical protein from BCRA2 region (CG005), mRNA
NM_014207	Homo sapiens CD5 antigen (p56-62) (CD5), mRNA
NM_014335	Homo sapiens chromosome 15 open reading frame 3 (C15ORF3), mRNA
NM_014206	Homo sapiens chromosome 11 open reading frame 10 (C11orf10), mRNA
NM_014453	Homo sapiens putative breast adenocarcinoma marker (32kD) (BC-2), mRNA
NM_014382	Homo sapiens ATPase, Ca++ transporting, type 2C, member 1 (ATP2C1), mRNA
NM_014570	Homo sapiens ADP-ribosylation factor GTPase activating protein 1 (ARFGAP1), mRNA
NM 014278	Homo sapiens heat shock protein (hsp110 family) (APG-1), mRNA
NM 014495	Homo sapiens angiopoietin-like 3 (ANGPTL3), mRNA
NM_004037	Homo sapiens adenosine monophosphate deaminase 2 (isoform L) (AMPD2), mRNA
NM_014324	Homo sapiens alpha-methylacyl-CoA racemase (AMACR), mRNA
NM 014476	Homo sapiens alpha-actinin-2-associated LIM protein (ALP), mRNA
NM 014423	Homo sapiens ALL1 fused gene from 5q31 (AF5Q31), mRNA
NM_014590	Homo sapiens endogenous retroviral family W, env(C7), member 1 (syncytin) (ERVWE1), mRNA
NM_014486	Homo sapiens neuronal thread protein (AD7C-NTP), mRNA
NM_014384	Homo sapiens acyl-Coenzyme A dehydrogenase family, member 8 (ACAD8), mRNA
NM_014274	Homo sapiens Alu-binding protein with zinc finger domain (ABP/ZF), mRNA
NM 014444	Homo sapiens gamma tubulin ring complex protein (76p gene) (76P), mRNA
NM_007082	Homo sapiens RAB, member of RAS oncogene family-like 2A (RABL2A), mRNA
NM_013412	Homo sapiens RAB, member of RAS oncogene family-like 2A (RABL2A), transcript variant 1, mRNA
NM_005036	Homo sapiens peroxisome proliferative activated receptor, alpha (PPARA), mRNA
NM_000793	Homo sapiens deiodinase, iodothyronine, type II (DIO2), transcript variant 2, mRNA
NM_013989	Homo sapiens deiodinase, iodothyronine, type II (DIO2), transcript variant 1, mRNA

NM_004323	Homo sapiens BCL2-associated athanogene (BAG1), mRNA
NM 000156	Homo sapiens guanidinoacetate N-methyltransferase (GAMT), mRNA
NM 002782	Homo sapiens pregnancy specific beta-1-glycoprotein 6 (PSG6), mRNA
NM 005523	Homo sapiens homeo box A11 (HOXA11), mRNA
NM 007050	Homo sapiens protein tyrosine phosphatase, receptor type, T (PTPRT), mRNA
NM 006249	Homo sapiens proline-rich protein BstNI subfamily 3 (PRB3), mRNA
NM 005529	Homo sapiens heparan sulfate proteoglycan 2 (perlecan) (HSPG2), mRNA
NM 005187	Homo sapiens core-binding factor, runt domain, alpha subunit 2; translocated to,
	3 (CBFA2T3), mRNA
NM_005565	Homo sapiens lymphocyte cytosolic protein 2 (SH2 domain-containing
	leukocyte protein of 76kD) (LCP2), mRNA
NM_002298	Homo sapiens lymphocyte cytosolic protein 1 (L-plastin) (LCP1), mRNA
NM 005190	Homo sapiens cyclin C (CCNC), mRNA
NM_005415	Homo sapiens solute carrier family 20 (phosphate transporter), member 1 (SLC20A1), mRNA
NM_001040	Homo sapiens sex hormone-binding globulin (SHBG), mRNA
	Homo sapiens proteinase 3 (serine proteinase, neutrophil, Wegener
NM_002777	granulomatosis autoantigen) (PRTN3), mRNA
NM_005199	Homo sapiens cholinergic receptor, nicotinic, gamma polypeptide (CHRNG),
NIVI_003199	mRNA
NM_013936	Homo sapiens olfactory receptor, family 12, subfamily D, member 2 (OR12D2), mRNA
NM 013937	Homo sapiens olfactory receptor, family 11, subfamily A, member 1 (OR11A1),
-	mRNA
NM_013940	Homo sapiens olfactory receptor, family 10, subfamily H, member 1 (OR10H1), mRNA
NM_013941	Homo sapiens olfactory receptor, family 10, subfamily C, member 1 (OR10C1), mRNA
NM_013938	Homo sapiens olfactory receptor, family 10, subfamily H, member 3 (OR10H3), mRNA
NM_013939	Homo sapiens olfactory receptor, family 10, subfamily H, member 2 (OR10H2), mRNA
NM 013452	Homo sapiens variable charge, X chromosome (VCX), mRNA
NM 013437	Homo sapiens potential tumor suppressor (ST7), mRNA
NM 013440	Homo sapiens paired immunoglobulin-like receptor beta (PILR(BETA)), mRNA
NM_013439	Homo sapiens paired immunoglobulin-like receptor alpha (PILR(ALPHA)), mRNA
NM 013446	Homo sapiens makorin, ring finger protein, 1 (MKRN1), mRNA
NM_007267	Homo sapiens expressed in activated T/LAK lymphocytes (LAK-4P), mRNA
NM 013450	Homo sapiens bromodomain adjacent to zinc finger domain, 2B (BAZ2B),
14141_013420	mRNA
NM_013448	Homo sapiens bromodomain adjacent to zinc finger domain, 1A (BAZ1A),
	mRNA
NM 000033	Homo sapiens ATP-binding cassette, sub-family D (ALD), member 1 (ABCD1),
	mRNA
NM 002593	Homo sapiens procollagen C-endopeptidase enhancer (PCOLCE), mRNA
NM_004504	Homo sapiens HIV-1 Rev binding protein (HRB), mRNA
NM_004131	Homo sapiens granzyme B (granzyme 2, cytotoxic T-lymphocyte-associated
1,11,1-10,13,1	serine esterase 1) (GZMB), mRNA
NM 000791	Homo sapiens dihydrofolate reductase (DHFR), mRNA
NM 004335	Homo sapiens bone marrow stromal cell antigen 2 (BST2), mRNA
NM 001197	Homo sapiens BCL2-interacting killer (apoptosis-inducing) (BIK), mRNA
TATAT OOT 12/	Liouno sabiens DCF7-interacting wifer (abobiosis-inducing) (DTG), interacting

NM_000487	Homo sapiens arylsulfatase A (ARSA), mRNA
NM_004597	Homo sapiens small nuclear ribonucleoprotein D2 polypeptide (16.5kD)
-	(SNRPD2), mRNA
NM_006194	Homo sapi ns paired box gene 9 (PAX9), mRNA
NM 013330	Homo sapiens NME7 (NME7), mRNA
NM_012476	Homo sapiens ventral anterior homeobox 2 (VAX2), mRNA
NM 012253	Homo sapiens transketolase-like 1 (TKTL1), mRNA
NM 012268	Homo sapiens similar to vaccinia virus HindIII K4L ORF (HU-K4), mRNA
NM 002017	Homo sapiens Friend leukemia virus integration 1 (FLI1), mRNA
NM 006769	Homo sapiens LIM domain only 4 (LMO4), mRNA
NM 002260	Homo sapiens killer cell lectin-like receptor subfamily C, member 2 (KLRC2),
-	mRNA
NM_005317	Homo sapiens granzyme M (lymphocyte met-ase 1) (GZMM), mRNA
NM 004417	Homo sapiens dual specificity phosphatase 1 (DUSP1), mRNA
NM 012125	Homo sapiens cholinergic receptor, muscarinic 5 (CHRM5), mRNA
NM 001236	Homo sapiens carbonyl reductase 3 (CBR3), mRNA
NM 013343	Homo sapiens NAG-7 protein (NAG-7), mRNA
NM_013344	Homo sapiens leucine zipper-like protein (LZLP), mRNA
NM 013236	Homo sapiens like mouse brain protein E46 (E46L), mRNA
NM 013380	Homo sapiens zinc finger protein 228 (ZNF228), mRNA
NM 013362	Homo sapiens zinc finger protein 225 (ZNF225), mRNA
NM 013398	Homo sapiens zinc finger protein 224 (ZNF224), mRNA
NM 013361	Homo sapiens zinc finger protein 223 (ZNF223), mRNA
NM 013360	Homo sapiens zinc finger protein 222 (ZNF222), mRNA
NM 013359	Homo sapiens zinc finger protein 221 (ZNF221), mRNA
NM 013250	Homo sapiens zinc finger protein 215 (ZNF215), mRNA
NM 013249	Homo sapiens zinc finger protein 214 (ZNF214), mRNA
NM 013256	Homo sapiens zinc finger protein 180 (HHZ168) (ZNF180), mRNA
NM 013371	Homo sapiens interleukin 19 (IL19), mRNA
NM 013403	Homo sapiens zinedin (ZIN), mRNA
NM 013378	Homo sapiens pre-B lymphocyte gene 3 (VPREB3), mRNA
NM 013270	Homo sapiens testes-specific protease 50 (TSP50), mRNA
NM_013381	Homo sapiens thyrotropin-releasing hormone degrading ectoenzyme (TRHDE),
	mRNA
NM 013315	Homo sapiens transmembrane phosphatase with tensin homology (TPTE),
-	mRNA
NM 013353	Homo sapiens tropomodulin 4 (muscle) (TMOD4), mRNA
NM_013390	Homo sapiens transmembrane protein 2 (TMEM2), mRNA
NM_013319	Homo sapiens transitional epithelia response protein (TERE1), mRNA
NM 013254	Homo sapiens TANK-binding kinase 1 (TBK1), mRNA
NM_013309	Homo sapiens solute carrier family 30 (zinc transporter), member 4 (SLC30A4),
İ	mRNA
NM_013356	Homo sapiens monocarboxylate transporter 3 (SLC16A8), mRNA
NM_013257	Homo sapiens serum/glucocorticoid regulated kinase-like (SGKL), mRNA
NM_013376	Homo sapiens CDK4-binding protein p34SEI1 (SEI1), mRNA
NM 013243	Homo sapiens secretogranin III (SCG3), mRNA
NM_013352	Homo sapiens squamous cell carcinoma antigen recognized by T cell (SART-2), mRNA
NM 013401	Homo sapiens RAB3A interacting protein (rabin3)-like 1 (RAB3IL1), mRNA
NM 013237	Homo sapiens px19-like protein (PX19), mRNA
NM_013261	Homo sapiens peroxisome proliferative activated receptor, gamma, coactivator 1
	(PPARGC1), mRNA

NM_013268	Homo sapiens placental protein 13 (PP13), mRNA
NM_013382	Homo sapiens putative protein O-mannosyltransferase (POMT2), mRNA
NM_013232	Homo sapiens programmed cell death 6 (PDCD6), mRNA
NM 013397	Homo sapiens over-expressed breast tumor protein (OBTP), mRNA
NM_013389	Homo sapiens NPC1 (Niemann-Pick disease, type C1, gene)-like 1 (NPC1L1), mRNA
NM 013326	Homo sapiens colon cancer-associated protein Mic1 (MIC1), mRNA
NM 013238	Homo sapiens DNAJ domain-containing (MCJ), mRNA
NM 013269	Homo sapiens lectin-like NK cell receptor (LLT1), mRNA
NM_013289	Homo sapiens killer cell immunoglobulin-like receptor, three domains, long cytoplasmic tail, 1 (KIR3DL1), mRNA
NM_013311	Homo sapiens insulin upstream factor 1 (IUF1), mRNA
NM 013278	Homo sapiens interleukin 17C (IL17C), mRNA
NM_013292	Homo sapiens (clone PWHLC2-24) myosin light chain 2 (HUMMLC2B), mRNA
NM_013288	Homo sapiens DNA binding protein for surfactant protein B (HUMBINDC), mRNA
NM_013244	Homo sapiens UDP-N-acetylglucosamine:a-1,3-D-mannoside beta-1,4-N-acetylglucosaminyltransferase IV-homolog (HGNT-IV-H), mRNA
NM 013264	Homo sapiens gonadotropin-regulated testicular RNA helicase (GRTH), mRNA
NM_013281	Homo sapiens fibronectin leucine rich transmembrane protein 3 (FLRT3), mRNA
NM_013231	Homo sapiens fibronectin leucine rich transmembrane protein 2 (FLRT2), mRNA
NM 013241	Homo sapiens FH1/FH2 domain-containing protein (FHOS), mRNA
NM_013342	Homo sapiens TCF3 (E2A) fusion partner (in childhood Leukemia) (TFPT), mRNA
NM_013246	Homo sapiens cardiotrophin-like cytokine; neurotrophin-1/B-cell stimulating factor-3 (CLC), mRNA
NM_013372	Homo sapiens cysteine knot superfamily 1, BMP antagonist 1 (CKTSF1B1), mRNA
NM 013327	Homo sapiens CGI-56 protein (CGI-56), mRNA
NM_013230	Homo sapiens CD24 antigen (small cell lung carcinoma cluster 4 antigen) (CD24), mRNA
NM 013276	Homo sapiens carbohydrate kinase-like (CARKL), mRNA
NM 013399	Homo sapiens chromosome 16 open reading frame 5 (C16orf5), mRNA
NM 006765	Homo sapiens Putative prostate cancer tumor suppressor (N33), mRNA
NM 006792	Homo sapiens mortality factor 4 (MORF4), mRNA
NM_000397	Homo sapiens cytochrome b-245, beta polypeptide (chronic granulomatous disease) (CYBB), mRNA
NM 005098	Homo sapiens musculin (activated B-cell factor-1) (MSC), mRNA
NM_006144	Homo sapiens granzyme A (granzyme 1, cytotoxic T-lymphocyte-associated serine esterase 3) (GZMA), mRNA
NM 002047	Homo sapiens glycyl-tRNA synthetase (GARS), mRNA
NM 004405	Homo sapiens distal-less homeo box 2 (DLX2), mRNA
NM 004371	Homo sapiens coatomer protein complex, subunit alpha (COPA), mRNA
NM_005181	Homo sapiens carbonic anhydrase III, muscle specific (CA3), mRNA
NM_001663	Homo sapiens ADP-ribosylation factor 6 (ARF6), mRNA
NM 001662	Homo sapiens ADP-ribosylation factor 5 (ARF5), mRNA
NM 001660	Homo sapiens ADP-ribosylation factor 4 (ARF4), mRNA
NM 001658	Homo sapiens ADP-ribosylation factor 1 (ARF1), mRNA
	Homo sapiens Cystic fibrosis transmembrane conductance regulator, ATP-
NM_000492	I Homo sapiens cysuc morosis transmemorane conductance regulator, ATT

	Line (1 5 - 1) Complete 7) (CINED) DAIA
	binding cassette (sub-family C, member 7) (CFTR), mRNA
NM_003560	Homo sapiens phospholipase A2, group VI (cytosolic, calcium-independent)
	(PLA2G6), mRNA
NM_004004	Homo sapiens gap junction protein, beta 2, 26kD (connexin 26) (GJB2), mRNA
NM_005198	Homo sapiens choline kinase-like (CHKL), mRNA
NM 012482	Homo sapiens zinc finger protein 281 (ZNF281), mRNA
NM 012256	Homo sapiens zinc finger protein 212 (ZNF212), mRNA
NM 012479	Homo sapiens tyrosine 3-monooxygenase/tryptophan 5-monooxygenase
_	activation protein, gamma polypeptide (YWHAG), mRNA
NM_012255	Homo sapiens 5'-3' exoribonuclease 2 (XRN2), mRNA
NM 012474	Homo sapiens uridine monophosphate kinase (UMPK), mRNA
NM 012473	Homo sapiens thioredoxin, mitochondrial (TXN2), mRNA
NM 012466	Homo sapiens tetraspanin TM4-B (TM4-B), mRNA
NM 012465	Homo sapiens tolloid-like 2 (TLL2), mRNA
NM 012464	Homo sapiens tolloid-like 1 (TLL1), mRNA
NM 012290	Homo sapiens tousled-like kinase 1 (TLK1), mRNA
NM 012455	Homo sapiens SEC7 homolog (TIC), mRNA
NM 012454	Homo sapiens T-cell lymphoma invasion and metastasis 2 (TIAM2), mRNA
	Homo sapiens transcription factor A, mitochondrial (TFAM), mRNA
NM_012251	Homo sapiens synaptogyrin 4 (SYNGR4), mRNA
NM_012451	Homo sapiens synaphogym 4 (STACK4), mactri Homo sapiens signal transducer and activator of transcription 5B (STAT5B),
NM_012448	mRNA
NM 012447	Homo sapiens stromal antigen 3 (STAG3), mRNA
NM 012445	Homo sapiens spondin 2, extracellular matrix protein (SPON2), mRNA
NM 012443	Homo sapiens sperm associated antigen 6 (SPAG6), mRNA
NM 012244	Homo sapiens solute carrier family 7 (cationic amino acid transporter, y+
	system), member 8 (SLC7A8), mRNA
NM_012243	Homo sapiens solute carrier family 35 (UDP-N-acetylglucosamine (UDP-
	GlcNAc) transporter), member 3 (SLC35A3), mRNA
NM 012434	Homo sapiens solute carrier family 17 (anion/sugar transporter), member 5
	(SLC17Å5), mRNA
NM 012432	Homo sapiens SET domain, bifurcated 1 (SETDB1), mRNA
NM 012427	Homo sapiens kallikrein 5 (KLK5), mRNA
NM 012236	Homo sapiens sex comb on midleg homolog 1 (SCMH1), mRNA
NM_012424	Homo sapiens ribosomal protein S6 kinase, 52kD, polypeptide 1 (RPS6KC1),
	mRNA
NM 012421	Homo sapiens rearranged L-myc fusion sequence (RLF), mRNA
NM_012415	Homo sapiens RAD54, S. cerevisiae, homolog of, B (RAD54B), mRNA
NM 012410	Homo sapiens type I transmembrane receptor (seizure-related protein) (PSK-1),
1,11,1_0,12,110	mRNA
NM 012409	Homo sapiens prion gene complex, downstream (PRND), mRNA
NM 012402	Homo sapiens partner of RAC1 (arfaptin 2) (POR1), mRNA
NM 012402	Homo sapiens phospholipase A2, group IID (PLA2G2D), mRNA
NM_012399	Homo sapiens phosphotidylinositol transfer protein, beta (PITPNB), mRNA
	Homo sapiens 6-phosphogluconolactonase (PGLS), mRNA
NM_012088	Homo sapiens o-phosphoglucoholactonase (FGES), hindva Homo sapiens PFTAIRE protein kinase 1 (PFTK1), mRNA
NM 012395	Homo sapiens PFTAIRE protein kinase 1 (FFTK1), inktya Homo sapiens prostate epithelium-specific Ets transcription factor (PDEF),
NM_012391	mRNA
NM_012385	Homo sapiens p8 protein (candidate of metastasis 1) (P8), mRNA
NM_012383	Homo sapiens osteoclast stimulating factor 1 (OSTF1), mRNA
NM 012375	Homo sapiens olfactory receptor, family 52, subfamily A, member 1 (OR52A1),
_	mRNA

NM_012368	Homo sapiens olfactory receptor, family 2, subfamily C, member 1 (OR2C1), mRNA
NM_012360	Homo sapiens olfactory receptor, family 1, subfamily F, member 8 (OR1F8), mRNA
NM_012352	Homo sapiens olfactory receptor, family 1, subfamily A, member 2 (OR1A2), mRNA
NM_012351	Homo sapiens olfactory receptor, family 10, subfamily J, member 1 (OR10J1), mRNA
NM_012345	Homo sapiens nuclear fragile X mental retardation protein interacting protein 1 (NUFIP1), mRNA
NM 012344	Homo sapiens neurotensin receptor 2 (NTSR2), mRNA
NM 012343	Homo sapiens nicotinamide nucleotide transhydrogenase (NNT), mRNA
NM 012342	Homo sapiens putative transmembrane protein (NMA), mRNA
NM 012337	Homo sapiens nasopharyngeal epithelium specific protein 1 (NESG1), mRNA
NM 012330	Homo sapiens histone acetyltransferase (MORF), mRNA
NM 012064	Homo sapiens major intrinsic protein of lens fiber (MIP), mRNA
NM 012214	Homo sapiens mannosyl (alpha-1,3-)-glycoprotein beta-1,4-N-
1111_011	acetylglucosaminyltransferase, isoenzyme A (MGAT4A), mRNA
NM 012213	Homo sapiens malonyl-CoA decarboxylase (MLYCD), mRNA
NM 012325	Homo sapiens microtubule-associated protein, RP/EB family, member 1
_	(MAPREI), mRNA
NM_012318	Homo sapiens leucine zipper-EF-hand containing transmembrane protein 1
	(LETM1), mRNA
NM_012317	Homo sapiens leucine zipper, down-regulated in cancer 1 (LDOC1), mRNA
NM_012314	Homo sapiens killer cell immunoglobulin-like receptor, two domains, short cytoplasmic tail, 4 (KIR2DS4), mRNA
NM_012313	Homo sapiens killer cell immunoglobulin-like receptor, two domains, short cytoplasmic tail, 3 (KIR2DS3), mRNA
NM_012312	Homo sapiens killer cell immunoglobulin-like receptor, two domains, short cytoplasmic tail, 2 (KIR2DS2), mRNA
NM_012307	Homo sapiens differentially expressed in adenocarcinoma of the lung (KIAA0987), mRNA
NM 012306	Homo sapiens lifeguard (KIAA0950), mRNA
NM 012302	Homo sapiens latrophilin (KIAA0786), mRNA
NM 012295	Homo sapiens calcineurin binding protein 1 (KIAA0330), mRNA
NM 012288	Homo sapiens TRAM-like protein (KIAA0057), mRNA
NM 012286	Homo sapiens MORF-related gene X (KIAA0026), mRNA
NM_012283	Homo sapiens potassium voltage-gated channel, subfamily G, member 2 (KCNG2), mRNA
NM_012282	Homo sapiens potassium voltage-gated channel, Isk-related family, member 1-like (KCNE1L), mRNA
NM 012278	Homo sapiens integrin beta 1 binding protein (melusin) 2 (ITGB1BP2), mRNA
NM 012211	Homo sapiens integrin, alpha 11 (ITGA11), mRNA
NM 012277	Homo sapiens pancreatic beta cell growth factor (INGAP), mRNA
NM 012277	Homo sapiens interleukin-1 receptor antagonist homolog 1 (IL1HY1), mRNA
NM_012259	Homo sapiens hairy/enhancer-of-split related with YRPW motif 2 (HEY2), mRNA
NM_012258	Homo sapiens hairy/enhancer-of-split related with YRPW motif 1 (HEY1), mRNA
NM_012257	Homo sapiens HMG-box containing protein 1 (HBP1), mRNA
NM 012087	Homo sapiens general transcription factor IIIC, polypeptide 5 (63kD) (GTF3C5),
I NIM DIZON	

	(CDYIDD)
NM_012203	Homo sapiens glyoxylate reductase/hydroxypyruvate reductase (GRHPR), mRNA
NM_012202	Homo sapiens guanine nucleotide binding protein (G protein), gamma 3 (GNG3), mRNA
NM_012084	Homo sapiens Glutamate dehydrogenase-2 (GLUD2), mRNA
NM 012191	Homo sapiens putative tumor suppressor (FUS2), mRNA
NM 012185	Homo sapiens forkhead box E2 (FOXE2), mRNA
NM 012183	Homo sapiens forkhead box D3 (FOXD3), mRNA
NM 012153	Homo sapiens Ets homologous factor (EHF), mRNA
NM_012080	Homo sapiens DNA segment, numerous copies, expressed probes (GS1 gene) (DXF68S1E), mRNA
NM 012148	Homo sapiens double homeobox, 3 (DUX3), mRNA
NM 012147	Homo sapiens double homeobox, 2 (DUX2), mRNA
NM 012145	Homo sapiens deoxythymidylate kinase (thymidylate kinase) (DTYMK), mRNA
NM 012144	Homo sapiens dynein, axonemal, intermediate polypeptide, 1 (DNAI1), mRNA
NM_012140	Homo sapiens solute carrier family 25 (mitochondrial carrier; dicarboxylate transporter), member 10 (SLC25A10), mRNA
NM 012137	Homo sapiens dimethylarginine dimethylaminohydrolase 1 (DDAH1), mRNA
NM 012134	Homo sapiens leiomodin 1 (smooth muscle) (LMOD1), mRNA
NM 012133	Homo sapiens coatomer protein complex, subunit gamma 2 (COPG2), mRNA
NM 012132	Homo sapiens claudin 8 (CLDN8), mRNA
NM 012131	Homo sapiens claudin 17 (CLDN17), mRNA
NM 012130	Homo sapiens claudin 14 (CLDN14), mRNA
NM 012129	Homo sapiens claudin 12 (CLDN12), mRNA
NM 012127	Homo sapiens Cip1-interacting zinc finger protein (CIZ1), mRNA
NM_012126	Homo sapiens carbohydrate (N-acetylglucosamine 6-O) sulfotransferase 5 (CHST5), mRNA
NM_012075	Homo sapiens Conserved gene telomeric to alpha globin cluster (CGTHBA), mRNA
NM_012122	Homo sapiens carboxylesterase 3 (brain) (CES3), mRNA
NM_012116	Homo sapiens Cas-Br-M (murine) ectropic retroviral transforming sequence c (CBLC), mRNA
NM 012113	Homo sapiens carbonic anhydrase XIV (CA14), mRNA
NM 012071	Homo sapiens BUP protein (BUP), mRNA
NM 012110	Homo sapiens cystein-rich hydrophobic domain 2 (CHIC2), mRNA
NM 012109	Homo sapiens brain-specific membrane-anchored protein (BSMAP), mRNA
NM_012107	Homo sapiens bromodomain containing protein 75 kDa human homolog (BP75), mRNA
NM_012104	Homo sapiens beta-site APP-cleaving enzyme (BACE), mRNA
NM_012105	Homo sapiens beta-site APP-cleaving enzyme 2 (BACE2), mRNA
NM 012103	Homo sapiens ancient ubiquitous protein 1 (AUP1), mRNA
NM 012102	Homo sapiens arginine-glutamic acid dipeptide (RE) repeats (RERE), mRNA
NM_012099	Homo sapiens CD3-epsilon-associated protein; antisense to ERCC-1 (ASE-1), mRNA
NM 012098	Homo sapiens angiopoietin-like 2 (ANGPTL2), mRNA
NM_012067	Homo sapiens aldo-keto reductase family 7, member A3 (aflatoxin aldehyde reductase) (AKR7A3), mRNA
NM 012093	Homo sapiens adenylate kinase 5 (AK5), mRNA
NM_012066	Homo sapiens hypothetical protein (20D7-FC4), mRNA
NM 006276	Homo sapiens splicing factor, arginine/serine-rich 7 (35kD) (SFRS7), mRNA
NM 007054	Homo sapiens kinesin family member 3A (KIF3A), mRNA
NM 002201	Homo sapiens interferon stimulated gene (20kD) (ISG20), mRNA
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NIM 007274	Homo sapiens cytosolic acyl coenzyme A thioester hydrolase (HBACH), mRNA
NM_007274	Homo sapiens solute carrier family 9 (sodium/hydrogen exchanger), isoform 3
NM_004174	(SLC9A3), mRNA
NM 004525	Homo sapiens low density lipoprotein-related protein 2 (LRP2), mRNA
NM 003129	Homo sapiens squalene epoxidase (SQLE), mRNA
NM 003628	Homo sapiens plakophilin 4 (PKP4), mRNA
NM 003028	Homo sapiens amine oxidase, copper containing 3 (vascular adhesion protein 1)
	(AOC3), mRNA
NM_003322	Homo sapiens tubby like protein 1 (TULP1), mRNA
NM_002747	Homo sapiens mitogen-activated protein kinase 4 (MAPK4), mRNA
NM_002078	Homo sapiens golgi autoantigen, golgin subfamily a, 4 (GOLGA4), mRNA
NM_006421	Homo sapiens brefeldin A-inhibited guanine nucleotide-exchange protein 1 (BIG1), mRNA
NM 004282	Homo sapiens BCL2-associated athanogene 2 (BAG2), mRNA
NM_004304	Homo sapiens anaplastic lymphoma kinase (Ki-1) (ALK), mRNA
NM_001626	Homo sapiens v-akt murine thymoma viral oncogene homolog 2 (AKT2), mRNA
NM 000686	Homo sapiens angiotensin receptor 2 (AGTR2), mRNA
NM_006287	Homo sapiens tissue factor pathway inhibitor (lipoprotein-associated coagulation inhibitor) (TFPI), mRNA
NM_000944	Homo sapiens protein phosphatase 3 (formerly 2B), catalytic subunit, alpha isoform (calcineurin A alpha) (PPP3CA), mRNA
NM_001142	Homo sapiens amelogenin (X chromosome, amelogenesis imperfecta 1) (AMELX), mRNA
NM_001171	Homo sapiens ATP-binding cassette, sub-family C (CFTR/MRP), member 6 (ABCC6), mRNA
NM 007351	Homo sapiens multimerin (MMRN), mRNA
NM 007355	Homo sapiens heat shock 90kD protein 1, beta (HSPCB), mRNA
NM 007354	Homo sapiens putative GR6 protein (GR6), mRNA
NM_007353	Homo sapiens guanine nucleotide binding protein (G protein) alpha 12 (GNA12), mRNA
NM 007366	Homo sapiens phospholipase A2 receptor 1, 180kD (PLA2R1), mRNA
NM_007350	Homo sapiens pleckstrin homology-like domain, family A, member 1 (PHLDA1), mRNA
NM 007364	Homo sapiens integral type I protein (P24B), mRNA
NM 007342	Homo sapiens nucleoporin-like protein 1 (NLP_1), mRNA
NM 007361	Homo sapiens nidogen 2 (NID2), mRNA
NM_007341	Homo sapiens SH3 domain binding glutamic acid-rich protein (SH3BGR), mRNA
NM 007370	Homo sapiens replication factor C (activator 1) 5 (36.5kD) (RFC5), mRNA
NM_007348	Homo sapiens activating transcription factor 6 (ATF6), mRNA
NM_004850	Homo sapiens Rho-associated, coiled-coil containing protein kinase 2 (ROCK2), mRNA
NM_005574	Homo sapiens LIM domain only 2 (rhombotin-like 1) (LMO2), mRNA
NM 006094	Homo sapiens deleted in liver cancer 1 (DLC1), mRNA
NM 003658	Homo sapiens BarH-like homeobox 2 (BARX2), mRNA
NM 004209	Homo sapiens synaptogyrin 3 (SYNGR3), mRNA
NM 004879	Homo sapiens etoposide-induced mRNA (PIG8), mRNA
NM 005385	Homo sapiens natural killer-tumor recognition sequence (NKTR), mRNA
NM_005957	Homo sapiens 5,10-methylenetetrahydrofolate reductase (NADPH) (MTHFR), mRNA
NM 002248	Homo sapiens potassium intermediate/small conductance calcium-activated
	Tarana Ambadan basedan and and and and and and and and and

	channel, subfamily N, member 1 (KCNN1), mRNA
ND 6 001662	Homo sapiens interphotoreceptor matrix proteoglycan 1 (IMPG1), mRNA
NM_001563	Homo sapiens interphototeceptor matrix proteogrycan 1 (hvir-G1), mRNA Homo sapiens gap junction protein, alpha 5, 40kD (connexin 40) (GJA5), mRNA
NM_005266	Homo sapiens carboxypeptidase M (CPM), mRNA
NM_001874	Homo sapiens ankyrin-like with transmembrane domains 1 (ANKTM1), mRNA
NM_007332	
NM_003313	Homo sapiens tissue specific transplantation antigen P35B (TSTA3), mRNA Homo sapiens GDP dissociation inhibitor 2 (GDI2), mRNA
NM_001494	Homo sapiens GDF dissociation inhibitor 2 (GDI2), mRNA Homo sapiens acetyl-Coenzyme A acyltransferase 1 (peroxisomal 3-oxoacyl-
NM_001607	Coenzyme A thiolase) (ACAA1), nuclear gene encoding mitochondrial protein,
1	mRNA
NM 003145	Homo sapiens signal sequence receptor, beta (translocon-associated protein beta)
11111_005145	(SSR2), mRNA
NM 000852	Homo sapiens glutathione S-transferase pi (GSTP1), mRNA
NM 000827	Homo sapiens glutamate receptor, ionotropic, AMPA 1 (GRIA1), mRNA
NM 005252	Homo sapiens v-fos FBJ murine osteosarcoma viral oncogene homolog (FOS),
	mRNA
NM_005803	Homo sapiens flotillin 1 (FLOT1), mRNA
NM_004459	Homo sapiens fetal Alzheimer antigen (FALZ), mRNA
NM_004081	Homo sapiens deleted in azoospermia (DAZ), mRNA
NM_004055	Homo sapiens calpain 5 (CAPN5), mRNA
NM_004042	Homo sapiens arylsulfatase F (ARSF), mRNA
NM_003085	Homo sapiens synuclein, beta (SNCB), mRNA
NM_000612	Homo sapiens insulin-like growth factor 2 (somatomedin A) (IGF2), mRNA
NM_006995	Homo sapiens butyrophilin, subfamily 2, member A2 (BTN2A2), mRNA
NM_005739	Homo sapiens RAS guanyl releasing protein 1 (calcium and DAG-regulated)
	(RASGRP1), mRNA
NM_006267	Homo sapiens RAN binding protein 2 (RANBP2), mRNA
NM_002882	Homo sapiens RAN binding protein 1 (RANBP1), mRNA
NM_003884	Homo sapiens p300/CBP-associated factor (PCAF), mRNA
NM_005258	Homo sapiens GTP cyclohydrolase I feedback regulatory protein (GCHFR), mRNA
NM 001130	Homo sapiens amino-terminal enhancer of split (AES), mRNA
NM 001099	Homo sapiens acid phosphatase, prostate (ACPP), mRNA
NM 005155	Homo sapiens palmitoyl-protein thioesterase 2 (PPT2), mRNA
NM_006898	Homo sapiens homeo box D3 (HOXD3), mRNA
NM_006894	Homo sapiens flavin containing monooxygenase 3 (FMO3), mRNA
NM_004111	Homo sapiens flap structure-specific endonuclease 1 (FEN1), mRNA
NM_001828	Homo sapiens Charot-Leyden crystal protein (CLC), mRNA
NM_007315	Homo sapiens signal transducer and activator of transcription 1, 91kD (STAT1), mRNA
NM_005005	Homo sapiens NADH dehydrogenase (ubiquinone) 1 beta subcomplex, 9 (22kD, B22) (NDUFB9), mRNA
NM 003362	Homo sapiens uracil-DNA glycosylase (UNG), mRNA
NM 005221	Homo sapiens distal-less homeo box 5 (DLX5), mRNA
NM_000479	Homo sapiens anti-Mullerian hormone (AMH), mRNA
NM 005160	Homo sapiens adrenergic, beta, receptor kinase 2 (ADRBK2), mRNA
NM 001619	Homo sapiens adrenergic, beta, receptor kinase 1 (ADRBK1), mRNA
NM 001611	Homo sapiens acid phosphatase 5, tartrate resistant (ACP5), mRNA
NM_003403	Homo sapiens YY1 transcription factor (YY1), mRNA
NM_003793	Homo sapiens cathepsin F (CTSF), mRNA
NM_001922	Homo sapiens dopachrome tautomerase (dopachrome delta-isomerase, tyrosine-
	r lated protein 2) (DCT), mRNA

NM_006412	Homo sapiens 1-acylglycerol-3-phosphate O-acyltransferase 2 (lysophosphatidic acid acyltransferase, beta) (AGPAT2), mRNA
NM_000810	Homo sapiens gamma-aminobutyric acid (GABA) A receptor, alpha 5 (GABRA5), mRNA
ND 6 000420	Homo sapiens platelet-activating factor acetylhydrolase, isoform Ib, alpha
NM_000430	subunit (45kD) (PAFAH1B1), mRNA
NM_003006	Homo sapiens selectin P ligand (SELPLG), mRNA
NM 002634	Homo sapiens prohibitin (PHB), mRNA
NM_002410	Homo sapiens mannosyl (alpha-1,6-)-glycoprotein beta-1,6-N-acetyl-glucosaminyltransferase (MGAT5), mRNA
NM 002409	Homo sapiens mannosyl (beta-1,4-)-glycoprotein beta-1,4-N-
112.2_002.00	acetylglucosaminyltransferase (MGAT3), mRNA
NM_002408	Homo sapiens mannosyl (alpha-1,6-)-glycoprotein beta-1,2-N-
11112_002-100	acetylglucosaminyltransferase (MGAT2), mRNA
NM 002406	Homo sapiens mannosyl (alpha-1,3-)-glycoprotein beta-1,2-N-
14141_002400	acetylglucosaminyltransferase (MGAT1), mRNA
ND 4 005022	Homo sapiens mitogen-activated protein kinase kinase kinase 5 (MAP3K5),
NM_005923	mRNA
NM_002225	Homo sapiens isovaleryl Coenzyme A dehydrogenase (IVD), nuclear gene
	encoding mitochondrial protein, mRNA
NM_001480	Homo sapiens galanin receptor 1 (GALR1), mRNA
NM 001992	Homo sapiens coagulation factor II (thrombin) receptor (F2R), mRNA
NM 000677	Homo sapiens adenosine A3 receptor (ADORA3), mRNA
NM 002969	Homo sapiens mitogen-activated protein kinase 12 (MAPK12), mRNA
NM 001526	Homo sapiens hypocretin (orexin) receptor 2 (HCRTR2), mRNA
NM_003605	Homo sapiens O-linked N-acetylglucosamine (GlcNAc) transferase (UDP-N-
14141_005005	acetylglucosamine:polypeptide-N-acetylglucosaminyl transferase) (OGT), mRNA
NM_000885	Homo sapiens integrin, alpha 4 (antigen CD49D, alpha 4 subunit of VLA-4 receptor) (ITGA4), mRNA
NM_003197	Homo sapiens transcription elongation factor B (SIII), polypeptide 1-like (TCEB1L), mRNA
NM 006183	Homo sapiens neurotensin (NTS), mRNA
NM_002524	Homo sapiens neuroblastoma RAS viral (v-ras) oncogene homolog (NRAS), mRNA
NM 002478	Homo sapiens myogenic factor 3 (MYOD1), mRNA
NM_002475	Homo sapiens methylthioadenosine phosphorylase (MTAP), mRNA
NM 002436	Homo sapiens membrane protein, palmitoylated 1 (55kD) (MPP1), mRNA
NM 002377	Homo sapiens MAS1 oncogene (MAS1), mRNA
NM 002377	Homo sapiens lectin, galactoside-binding, soluble, 1 (galectin 1) (LGALS1),
	mRNA
NM_000887	Homo sapiens integrin, alpha X (antigen CD11C (p150), alpha polypeptide) (ITGAX), mRNA
NM_000419	Homo sapiens integrin, alpha 2b (platelet glycoprotein IIb of IIb/IIIa complex, antigen CD41B) (ITGA2B), mRNA
NM_002203	Homo sapiens integrin, alpha 2 (CD49B, alpha 2 subunit of VLA-2 receptor) (ITGA2), mRNA
NTM 002627	
NM_003637	Homo sapiens integrin, alpha 10 (ITGA10), mRNA
NM_000843	Homo sapiens glutamate receptor, metabotropic 6 (GRM6), mRNA
NM_000838	Homo sapiens glutamate receptor, metabotropic 1 (GRM1), mRNA
NM_000835	Homo sapiens glutamate receptor, ionotropic, N-methyl D-aspartate 2C (GRIN2C), mRNA

NM_000834	Homo sapiens glutamate receptor, ionotropic, N-methyl D-aspartate 2B (GRIN2B), mRNA
NM_000833	Homo sapiens glutamate receptor, ionotropic, N-methyl D-aspartate 2A
	(GRIN2A), mRNA
NM_002084	Homo sapiens glutathione peroxidase 3 (plasma) (GPX3), mRNA
NM_000805	Homo sapiens gastrin (GAS), mRNA
NM_001940	Homo sapiens dentatorubral-pallidoluysian atrophy (atrophin-1) (DRPLA), mRNA
NM_001219	Homo sapiens calumenin (CALU), mRNA
NM 007155	Homo sapiens zona pellucida glycoprotein 3A (sperm receptor) (ZP3A), mRNA
NM 007136	Homo sapiens zinc finger protein 80 (pT17) (ZNF80), mRNA
NM 007250	Homo sapiens Kruppel-like factor 8 (KLF8), mRNA
NM 007167	Homo sapiens zinc finger protein 258 (ZNF258), mRNA
NM 007153	Homo sapiens zinc finger protein 208 (ZNF208), mRNA
NM 007152	Homo sapiens zinc finger protein 195 (ZNF195), mRNA
NM 007150	Homo sapiens zinc finger protein 185 (LIM domain) (ZNF185), mRNA
NM 007147	Homo sapiens zinc finger protein 175 (ZNF175), mRNA
NM 007145	Homo sapiens zinc finger protein 146 (ZNF146), mRNA
NM 007127	Homo saniens villin 1 (VIL1), mRNA
NM_007125	Homo sapiens ubiquitously transcribed tetratricopeptide repeat gene, Y chromosome (UTY), mRNA
NM 007124	Homo sapiens utrophin (homologous to dystrophin) (UTRN), mRNA
NM 007122	Homo sapiens upstream transcription factor 1 (USF1), mRNA
NM_007120	Homo sapiens UDP glycosyltransferase 2 family, polypeptide B (UGT2B), mRNA
NM 007106	Homo sapiens ubiquitin-like 3 (UBL3), mRNA
NM 007118	Homo sapiens triple functional domain (PTPRF interacting) (TRIO), mRNA
NM 007117	Homo sapiens thyrotropin-releasing hormone (TRH), mRNA
NM_007218	Homo sapiens patched related protein translocated in renal cancer (TRC8), mRNA
NM 007233	Homo sapiens TP53 target gene 1 (TP53TG1), mRNA
NM 007114	Homo sapiens TATA element modulatory factor 1 (TMF1), mRNA
NM 007112	Homo sapiens thrombospondin 3 (THBS3), mRNA
NM 007111	Homo sapiens transcription factor Dp-1 (TFDP1), mRNA
NM 007109	Homo sapiens transcription factor 19 (SC1) (TCF19), mRNA
NM_007108	Homo sapiens transcription elongation factor B (SIII), polypeptide 2 (18kD, elongin B) (TCEB2), mRNA
NM_007105	Homo sapiens solute carrier family 22 (organic cation transporter), member 1-like antisense (SLC22A1LS), mRNA
NM_007163	Homo sapiens solute carrier family 14 (urea transporter), member 2 (SLC14A2), mRNA
NM 007101	Homo sapiens sarcosine dehydrogenase (SARDH), mRNA
NM 007165	Homo sapiens splicing factor 3a, subunit 2, 66kD (SF3A2), mRNA
NM 007252	Homo sapiens Retina-derived POU-domain factor-1 (RPF-1), mRNA
	Homo sapiens B-cell associated protein (REA), mRNA
NM_007273	Homo sapiens B-cen associated protein (REA), michay Homo sapiens polymerase (DNA directed) iota (POLI), mRNA
NM_007195	Homo sapiens potein tyrosine kinase 9-like (A6-related protein) (PTK9L),
NM_007284	mRNA
NM_007196	Homo sapiens kallikrein 8 (neuropsin/ovasin) (KLK8), mRNA
NM_007171	Homo sapiens protein-O-mannosyltransferase 1 (POMT1), mRNA
NM_007215	Homo sapiens polymerase (DNA directed), gamma 2, accessory subunit (POLG2), mRNA

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NM_007254	Homo sapiens polynucleotide kinase 3'-phosphatas (PNKP), mRNA
NM_007221	Homo sapiens polyamine-modulated factor 1 (PMF1), mRNA
NM_007183	Homo sapiens plakophilin 3 (PKP3), mRNA
NM_007169	Homo sapiens phosphatidylethanolamine N-methyltransferase (PEMT), mRNA
NM 007229	Homo sapiens protein kinase C and casein kinase substrate in neurons 2
_	(PACSIN2), mRNA
NM 007190	Homo sapiens Sec23-interacting protein p125 (P125), mRNA
NM 007160	Homo sapiens olfactory receptor, family 2, subfamily H, member 3 (OR2H3),
_	mRNA
NM 007256	Homo sapiens solute carrier family 21 (organic anion transporter), member 9
	(SLC21A9), mRNA
NM 007172	Homo sapiens nucleoporin 50kD (NUP50), mRNA
NM 007103	Homo sapiens NADH dehydrogenase (ubiquinone) flavoprotein 1 (51kD)
	(NDUFV1), mRNA
NM 007181	Homo sapiens mitogen-activated protein kinase kinase kinase 1
	(MAP4K1), mRNA
NM 007230	Homo sapiens mannosidase, alpha, class 1B, member 1 (MAN1B1), mRNA
NM_007164	Homo sapiens mucosal vascular addressin cell adhesion molecule 1
11112_007101	(MADCAM1), mRNA
NM 007216	Homo sapiens alpha integrin binding protein 63 (KIAA1017), mRNA
NM 007213	Homo sapiens JM4 protein (JM4), mRNA
NM_007102	Homo sapiens guanylate cyclase activator 2B (uroguanylin) (GUCA2B), mRNA
NM 007227	Homo sapiens G protein-coupled receptor 45 (GPR45), mRNA
NM_007275	Homo sapiens lung cancer candidate (FUS1), mRNA
NM_007262	Homo sapiens RNA-binding protein regulatory subunit (DJ-1), mRNA
NM 007166	Homo sapiens Clathrin assembly lymphoid-myeloid leukemia gene (CLTH),
14141_007100	mRNA
NM_007186	Homo sapiens centrosomal protein 2 (CEP2), mRNA
NM 006585	Homo sapiens chaperonin containing TCP1, subunit 8 (theta) (CCT8), mRNA
NM 007185	Homo sapiens trinucleotide repeat containing 4 (TNRC4), mRNA
NM_007220	Homo sapiens carbonic anhydrase VB, mitochondrial (CA5B), nuclear gene
14141_007220	encoding mitochondrial protein, mRNA
NM_007100	Homo sapiens ATP synthase, H+ transporting, mitochondrial F0 complex,
14141_007100	subunit e (ATP5I), mRNA
NM_007231	Homo sapiens solute carrier family 6 (neurotransmitter transporter), member 14
1414_007251	(SLC6A14), mRNA
NM 007203	Homo sapiens A kinase (PRKA) anchor protein 2 (AKAP2), mRNA
NM 007202	Homo sapiens A kinase (PRKA) anchor protein 10 (AKAP10), mRNA
NM 007168	Homo sapiens ATP-binding cassette, sub-family A (ABC1), member 8
14141_00\109	(ABCA8), mRNA
NM_000506	Homo sapiens coagulation factor II (thrombin) (F2), mRNA
NM 004343	Homo sapiens coagulation factor if (missinom) (12), matter Homo sapiens calreticulin (CALR), mRNA
NM 006736	Homo sapiens teat shock protein, neuronal DNAJ-like 1 (HSJ1), mRNA
NM 006553	Homo sapiens erythroid differentiation and denucleation factor 1 (HFL-EDDG1),
14141 000222	mRNA
NM 006984	Homo sapiens claudin 10 (CLDN10), mRNA
	Homo sapiens ATP-binding cassette, sub-family A (ABC1), member 1
NM_005502	(ABCA1), mRNA
ND 6 005000	
NM_005809	Homo sapiens peroxiredoxin 2 (PRDX2), mRNA
NM_006977	Homo sapiens zinc finger protein 46 (KUP) (ZNF46), mRNA
NM_006965	Homo sapiens zinc finger protein 24 (KOX 17) (ZNF24), mRNA
NM_006963	Homo sapiens zinc finger protein 22 (KOX 15) (ZNF22), mRNA

NM_006978	Homo sapiens zinc finger protein 183 (RING finger, C3HC4 type) (ZNF183), mRNA
NM 006953	Homo sapiens uroplakin 3 (UPK3), mRNA
NM 006952	Homo sapiens uroplakin 1B (UPK1B), mRNA
NM_006951	Homo sapiens TATA box binding protein (TBP)-associated factor, RNA
	polymerase II, D, 100kD (TAF2D), mRNA
NM 006950	Homo sapiens synapsin I (SYN1), mRNA
NM 007056	Homo sapiens suppressor of white apricot homolog 2 (SWAP2), mRNA
NM 006949	Homo sapiens syntaxin binding protein 2 (STXBP2), mRNA
NM 006948	Homo sapiens stress 70 protein chaperone, microsome-associated, 60kD
	(STCH), mRNA
NM 006946	Homo sapiens spectrin, beta, non-erythrocytic 2 (SPTBN2), mRNA
NM 006945	Homo sapiens small proline-rich protein 2B (SPRR2B), mRNA
NM 006944	Homo sapiens secreted phosphoprotein 2, 24kD (SPP2), mRNA
NM 007009	Homo sapiens zona pellucida binding protein (SP38), mRNA
NM 006940	Homo sapiens SRY (sex determining region Y)-box 5 (SOX5), mRNA
NM 007017	Homo sapiens SRY (sex determining region Y)-box 30 (SOX30), mRNA
NM_006943	Homo sapiens SRY (sex determining region Y)-box 22 (SOX22), mRNA
NM_007084	Homo sapiens SRY (sex determining region Y)-box 21 (SOX21), mRNA
NM_006942	Homo sapiens SRY (sex determining region Y)-box 20 (SOX20), mRNA
NM 006941	Homo sapiens SRY (sex determining region Y)-box 10 (SOX10), mRNA
NM_006934	Homo sapiens solute carrier family 6 (neurotransmitter transporter, glycine),
	member 9 (SLC6A9), mRNA
NM_006933	Homo sapiens solute carrier family 5 (inositol transporters), member 3
_	(SLC5A3), mRNA
NM 006931	Homo sapiens solute carrier family 2 (facilitated glucose transporter), member 3
	(SLC2A3), mRNA
NM_006930	Homo sapiens S-phase kinase-associated protein 1A (p19A) (SKP1A), mRNA
NM_006925	Homo sapiens splicing factor, arginine/serine-rich 5 (SFRS5), mRNA
NM_006924	Homo sapiens splicing factor, arginine/serine-rich 1 (splicing factor 2, alternate
	splicing factor) (SFRS1), mRNA
NM_006917	Homo sapiens retinoid X receptor, gamma (RXRG), mRNA
NM_006987	Homo sapiens rabphilin 3A-like (without C2 domains) (RPH3AL), mRNA
NM_007055	Homo sapiens polymerase (RNA) III (DNA directed) (155kD) (RPC155),
	mRNA
NM_006915	Homo sapiens retinitis pigmentosa 2 (X-linked recessive) (RP2), mRNA
NM_006914	Homo sapiens RAR-related orphan receptor B (RORB), mRNA
NM_006913	Homo sapiens ring finger protein 5 (RNF5), mRNA
NM_006911	Homo sapiens relaxin 1 (H1) (RLN1), mRNA
NM_007043	Homo sapiens HIV-1 rev binding protein 2 (HRB2), mRNA
NM_007033	Homo sapiens similar to S. cerevisiae RER1 (RER1), mRNA
NM_007081	Homo sapiens RAB, member of RAS oncogene family-like 2B (RABL2B),
	mRNA PNA
NM_006905	Homo sapiens pregnancy specific beta-1-glycoprotein 1 (PSG1), mRNA
NM_007016	Homo sapiens protein similar to E.coli yhdg and R. capsulatus nifR3 (PP35), mRNA
NM_007024	Homo sapiens PL6 protein (PL6), mRNA
NM_007030	Homo sapiens brain-specific protein p25 alpha (p25), mRNA
NM_006901	Homo sapiens myosin IXA (MYO9A), mRNA
NM_007075	Homo sapiens JM5 protein (JM5), mRNA
NM_007003	Homo sapiens JM27 protein (JM27), mRNA
NM 006899	Homo sapiens isocitrate dehydrogenase 3 (NAD+) beta (IDH3B), mRNA

NM_007031	Homo sapiens heat shock transcription factor 2 binding protein (HSF2BP), mRNA
NM 007011	Homo sapiens putative transmembrane protein (HS1-2), mRNA
NM 006896	Homo sapiens homeo box A7 (HOXA7), mRNA
NM 007045	Homo sapiens FGFR1 oncogene partner (FOP), mRNA
NM 007051	Homo sapiens Fas (TNFRSF6) associated factor 1 (FAF1), mRNA
NM 006979	Homo sapiens HLA class II region expressed gene KE4 (HKE4), mRNA
NM 007015	Homo sapiens chondromodulin I precursor (CHM-I), mRNA
NM 006890	Homo sapiens carcinoembryonic antigen-related cell adhesion molecule 7
14141_000930	(CEACAM7), mRNA
NM_007018	Homo sapiens centrosomal protein 1 (CEP1), mRNA
NM_006889	Homo sapiens CD86 antigen (CD28 antigen ligand 2, B7-2 antigen) (CD86), mRNA
NM 006982	Homo sapiens cartilage paired-class homeoprotein 1 (CART1), mRNA
NM 007058	Homo sapiens calpain 11 (CAPN11), mRNA
NM 006888	Homo sapiens calmodulin 1 (phosphorylase kinase, delta) (CALM1), mRNA
NM 007047	Homo sapiens butyrophilin, subfamily 3, member A2 (BTN3A2), mRNA
NM 007048	Homo sapiens butyrophilin, subfamily 3, member A1 (BTN3A1), mRNA
NM 006992	Homo sapiens B7 protein (B7), mRNA
NM 006885	Homo sapiens AT-binding transcription factor 1 (ATBF1), mRNA
NM 007022	Homo sapiens putative tumor suppressor (101F6), mRNA
NM 006697	Homo sapiens cisplatin resistance associated (CRA), mRNA
NM 006826	Homo sapiens tyrosine 3-monooxygenase/tryptophan 5-monooxygenase
1444_000020	activation protein, theta polypeptide (YWHAQ), mRNA
NM_006761	Homo sapiens tyrosine 3-monooxygenase/tryptophan 5-monooxygenase
_	activation protein, epsilon polypeptide (YWHAE), mRNA
NM_006784	Homo sapiens WD repeat domain 3 (WDR3), mRNA
NM_006846	Homo sapiens serine protease inhibitor, Kazal type, 5 (SPINK5), mRNA
NM_006830	Homo sapiens ubiquinol-cytochrome c reductase (6.4kD) subunit (UQCR), mRNA
NM_006798	Homo sapiens UDP glycosyltransferase 2 family, polypeptide A1 (UGT2A1), mRNA
NM 006757	Homo sapiens troponin T3, skeletal, fast (TNNT3), mRNA
NM 006827	Homo sapiens transmembrane trafficking protein (TMP21), mRNA
NM 006853	Homo sapiens kallikrein 11 (KLK11), mRNA
NM 006811	Homo sapiens tumor differentially expressed 1 (TDE1), mRNA
NM 006756	Homo sapiens transcription elongation factor A (SII), 1 (TCEA1), mRNA
NM_006024	Homo sapiens Tax1 (human T-cell leukemia virus type I) binding protein 1
1111_000024	(TAX1BP1), mRNA
NM 006752	Homo sapiens surfeit 5 (SURF5), mRNA
NM 006819	Homo sapiens stress-induced-phosphoprotein 1 (Hsp70/Hsp90-organizing
1	protein) (STIP1), mRNA
NM_006780	Homo sapiens SMA3 (SMA3), mRNA
NM_006749	Homo sapiens solute carrier family 20 (phosphate transporter), member 2
	(SLC20A2), mRNA
NM_006747	Homo sapiens signal-induced proliferation-associated gene 1 (SIPA1), mRNA
NM 006873	Homo sapiens stoned B/TFIIA-alpha/beta-like factor (SALF), mRNA
NM 006788	Homo sapiens ralA binding protein 1 (RALBP1), mRNA
NM 006871	Homo sapiens receptor-interacting serine-threonine kinase 3 (RIPK3), mRNA
NM_006867	Homo sapiens RNA-binding protein gene with multiple splicing (RBPMS), mRNA
NM 006743	Homo sapiens RNA binding motif protein 3 (RBM3), mRNA
14147 000143	

NM 006868	Homo sapiens RAB31, member RAS oncogene family (RAB31), mRNA
NM_006839	Homo sapiens inner membrane protein, mitochondrial (mitofilin) (IMMT),
_	mRNA
NM 006812	Homo sapiens amplified in osteosarcoma (OS-9), mRNA
NM 006656	Homo sapiens sialidase 3 (membrane sialidase) (NEU3), mRNA
NM 006791	Homo sapiens MORF-related gene 15 (MRG15), mRNA
NM 006766	Homo sapiens zinc finger protein 220 (ZNF220), mRNA
NM 006804	Homo sapiens steroidogenic acute regulatory protein related (MLN64), mRNA
NM_006770	Homo sapiens macrophage receptor with collagenous structure (MARCO),
14141_000770	mRNA
NM_006785	Homo sapiens mucosa associated lymphoid tissue lymphoma translocation gene
MM_000193	1 (MALT1), mRNA
ND 6 006767	Homo sapiens leucine-zipper-like transcriptional regulator, 1 (LZTR1), mRNA
NM_006767	Homo sapiens leukocyte immunoglobulin-like receptor, subfamily B (with TM
NM_006840	and ITIM domains), member 5 (LILRB5), mRNA
	Homo sapiens leukocyte immunoglobulin-like receptor, subfamily A (with TM
NM_006866	Homo sapiens leukocyte immunoglobuliii-like tecepoli, suotainity A (with 1141
	domain), member 2 (LILRA2), mRNA
NM_006863	Homo sapiens leukocyte immunoglobulin-like receptor, subfamily A (with TM
	domain), member 1 (LILRA1), mRNA
NM_006847	Homo sapiens leukocyte immunoglobulin-like receptor, subfamily B (with TM
	and ITIM domains), member 4 (LILRB4), mRNA
NM_006865	Homo sapiens leukocyte immunoglobulin-like receptor, subfamily A (without
	TM domain), member 3 (LILRA3), mRNA
NM_006864	Homo sapiens leukocyte immunoglobulin-like receptor, subfamily B (with TM
	and ITIM domains), member 3 (LILRB3), mRNA
NM_006738	Homo sapiens lymphoid blast crisis oncogene (LBC), mRNA
NM_006762	Homo sapiens Lysosomal-associated multispanning membrane protein-5 (LAPTM5), mRNA
NM_006737	Homo sapiens killer cell immunoglobulin-like receptor, three domains, long
14141_000757	cytoplasmic tail, 2 (KIR3DL2), mRNA
NM 006801	Homo sapiens KDEL (Lys-Asp-Glu-Leu) endoplasmic reticulum protein
14141_000001	retention receptor 1 (KDELR1), mRNA
NM 006844	Homo sapiens ilvB (bacterial acetolactate synthase)-like (ILVBL), mRNA
	Homo sapiens putative T1/ST2 receptor binding protein (IL1RL1LG), mRNA
NM_006858	Homo sapiens interferon-related developmental regulator 2 (IFRD2), mRNA
NM_006764	Homo sapiens ATP/GTP-binding protein (HEAB), mRNA
NM 006831	Homo sapiens ATP/GTP-officing protein (HEAD), mixty
NM_006794	Homo sapiens G protein-coupled receptor 75 (GPR75), mRNA
NM_006783	Homo sapiens gap junction protein, beta 6 (connexin 30) (GJB6), mRNA
NM_006733	Homo sapiens FSH primary response (LRPR1, rat) homolog 1 (FSHPRH1), mRNA
NM 006731	Homo sapiens Fukuyama type congenital muscular dystrophy (FCMD), mRNA
NM_006730	Homo sapiens deoxyribonuclease I-like 1 (DNASE1L1), mRNA
NM 004366	Homo sapiens chloride channel 2 (CLCN2), mRNA
NM 006725	Homo sapiens CD6 antigen (CD6), mRNA
	Homo sapiens BTG family, member 3 (BTG3), mRNA
NM 006806	Homo sapiens BTG family, member 2 (BTG2), mRNA
NM_006763	Homo sapiens apolipoprotein B mRNA editing enzyme, catalytic polypeptide-
NM_006789	like 2 (APOBEC2), mRNA
NM_006793	Homo sapiens peroxiredoxin 3 (PRDX3), nuclear gene encoding mitochondrial
_ ,	protein, mRNA
NM_006818	Homo sapiens ALL1-fused gene from chromosome 1q (AF1Q), mRNA
	Homo sapiens nuclear factor (erythroid-derived 2)-like 3 (NFE2L3), mRNA

NIM ODCEDE	Homo sapiens zinc finger protein 217 (ZNF217), mRNA
NM_006526	Homo sapiens X-prolyl aminopeptidase (aminopeptidase P)-like (XPNPEPL),
NM_006523	mRNA
NM 006537	Homo sapiens ubiquitin specific protease 3 (USP3), mRNA
NM 006564	Homo sapiens G protein-coupled receptor (TYMSTR), mRNA
NM 006573	Homo sapiens tumor necrosis factor (ligand) superfamily, member 13b
112.7_000070	(TNFSF13B), mRNA
NM 001561	Homo sapiens tumor necrosis factor receptor superfamily, member 9
	(TNFRSF9), mRNA
NM 006528	Homo sapiens tissue factor pathway inhibitor 2 (TFPI2), mRNA
NM 006520	Homo sapiens t-complex-associated-testis-expressed 1-like (TCTE1L), mRNA
NM 006519	Homo sapiens t-complex-associated-testis-expressed 1-like 1 (TCTEL1), mRNA
NM 006602	Homo sapiens transcription factor-like 5 (basic helix-loop-helix) (TCFL5),
	mRNA
NM 006593	Homo sapiens T-box, brain, 1 (TBR1), mRNA
NM_006679	Homo sapiens putative opioid receptor, neuromedin K (neurokinin B) receptor-like (TAC3RL), mRNA
NM 006682	Homo sapiens fibrinogen-like 2 (FGL2), mRNA
NM 006558	Homo sapiens Sam68-like phosphotyrosine protein, T-STAR (T-STAR), mRNA
NM 006603	Homo sapiens stromal antigen 2 (STAG2), mRNA
NM 006717	Homo sapiens spindlin (SPIN), mRNA
NM 006542	Homo sapiens S-phase response (cyclin-related) (SPHAR), mRNA
NM 006654	Homo sapiens sucl-associated neurotrophic factor target (FGFR signalling
1411_000034	adaptor) (SNT-1), mRNA
NM_006622	Homo sapiens serum-inducible kinase (SNK), mRNA
NM_006696	Homo sapiens thyroid hormone receptor coactivating protein (SMAP), mRNA
NM_006516	Homo sapiens solute carrier family 2 (facilitated glucose transporter), member 1 (SLC2A1), mRNA
NM_006632	Homo sapiens solute carrier family 17 (sodium phosphate), member 3 (SLC17A3), mRNA
NM 006517	Homo sapiens solute carrier family 16 (monocarboxylic acid transporters),
_	member 2 (putative transporter) (SLC16A2), mRNA
NM_006598	Homo sapiens solute carrier family 12 (potassium/chloride transporters), member
	7 (SLC12A7), mRNA
NM_006515	Homo sapiens SET domain and mariner transposase fusion gene (SETMAR), mRNA
NM_006664	Homo sapiens small inducible cytokine subfamily A (Cys-Cys), member 27 (SCYA27), mRNA
NM_006514	Homo sapiens sodium channel, voltage-gated, type X, alpha polypeptide (SCN10A), mRNA
NM_006559	Homo sapiens GAP-associated tyrosine phosphoprotein p62 (Sam68) (SAM68), mRNA
NM_006511	Homo sapiens regulatory solute carrier protein, family 1, member 1 (RSC1A1), mRNA
NM_006583	Homo sapiens retinal pigment epithelium-derived rhodopsin homolog (RRH), mRNA
NM 006604	Homo sapiens ret finger protein-like 3 (RFPL3), mRNA
NM 006605	Homo sapiens ret finger protein-like 2 (RFPL2), mRNA
NM 006505	Homo sapiens poliovirus receptor (PVR), mRNA
NM 006504	Homo sapiens protein tyrosine phosphatase, receptor type, E (PTPRE), mRNA
NM_006503	Homo sapiens proteasome (prosome, macropain) 26S subunit, ATPase, 4
	(PSMC4), mRNA

NM_006587	Homo sapiens corin (PRSC), mRNA
NM_006556	Homo sapiens phosphomevalonate kinase (PMVK), mRNA
NM_006608	Homo sapiens putative homeodomain transcription factor (PHTF1), mRNA
NM_006661	Homo sapiens phosphodiesterase 10A (PDE10A), mRNA
NM_006674	Homo sapiens MHC class I region ORF (P5-1), mRNA
NM_006637	Homo sapiens olfactory receptor, family 5, subfamily I, member 1 (OR5I1), mRNA
NM_006649	Homo sapiens serologically defined colon cancer antigen 16 (SDCCAG16), mRNA
NM_002532	Homo sapiens nucleoporin 88kD (NUP88), mRNA
NM_006702	Homo sapiens neuropathy target esterase (NTE), mRNA
NM_006693	Homo sapiens cleavage and polyadenylation specific factor 4, 30kD subunit (CPSF4), mRNA
NM_006669	Homo sapiens leukocyte immunoglobulin-like receptor, subfamily B (with TM and ITIM domains), member 1 (LILRB1), mRNA
NM 006533	Homo sapiens melanoma inhibitory activity (MIA), mRNA
NM 006500	Homo sapiens melanoma adhesion molecule (MCAM), mRNA
NM 006610	Homo sapiens mannan-binding lectin serine protease 2 (MASP2), mRNA
NM 006699	Homo sapiens mannosidase, alpha, class 1A, member 2 (MAN1A2), mRNA
NM_006498	Homo sapiens lectin, galactoside-binding, soluble, 2 (galectin 2) (LGALS2), mRNA
NM_006547	Homo sapiens IGF-II mRNA-binding protein 3 (KOC1), mRNA
NM_006611	Homo sapiens killer cell lectin-like receptor subfamily A, member 1 (KLRA1), mRNA
NM_006546	Homo sapiens IGF-II mRNA-binding protein 1 (IMP-1), mRNA
NM 006665	Homo sapiens heparanase (HPSE), mRNA
NM 006497	Homo sapiens hypermethylated in cancer 1 (HIC1), mRNA
NM 004667	Homo sapiens hect domain and RLD 2 (HERC2), mRNA
NM 006527	Homo sapiens Hairpin binding protein, histone (HBP), mRNA
NM 006658	Homo sapiens G-substrate (GSBS), mRNA
NM_006496	Homo sapiens guanine nucleotide binding protein (G protein), alpha inhibiting
27.6.006500	activity polypeptide 3 (GNAI3), mRNA
NM_006529	Homo sapiens glycine receptor, alpha 3 (GLRA3), mRNA
NM_006530	Homo sapiens glioma-amplified sequence-41 (GAS41), mRNA Homo sapiens fucosyltransferase 9 (alpha (1,3) fucosyltransferase) (FUT9),
NM_006581	mRNA
NM 006700	Homo sapiens FLN29 gene product (FLN29), mRNA
NM_006684	Homo sapiens complement factor H-related 4 (FHR-4), mRNA
NM_004113	Homo sapiens fibroblast growth factor 12B (FGF12B), mRNA
NM_006495	Homo sapiens ecotropic viral integration site 2B (EVI2B), mRNA
NM_006532	Homo sapiens ELL gene (11-19 lysine-rich leukemia gene) (ELL), mRNA
NM_006566	Homo sapiens adhesion glycoprotein (DNAM-1), mRNA
NM 006639	Homo sapiens cysteinyl leukotriene receptor 1 (CYSLT1), mRNA
NM_006586	Homo sapiens trinucleotide repeat containing 5 (TNRC5), mRNA
NM_006565	Homo sapiens CCCTC-binding factor (zinc finger protein) (CTCF), mRNA
NM_006574	Homo sapiens chondroitin sulfate proteoglycan 5 (neuroglycan C) (CSPG5), mRNA
NM 006688	Homo sapiens C1q-related factor (CRF), mRNA
NM 006493	Homo sapiens ceroid-lipofuscinosis, neuronal 5 (CLN5), mRNA
NM 001750	Homo sapiens calpastatin (CAST), mRNA
NM 006624	Homo sapiens adenovirus 5 E1A binding protein (BS69), mRNA
NM 006698	Homo sapiens bladder cancer associated protein (BLCAP), mRNA
NM_006698	Homo sapiens bladder cancer associated protein (BLCAP), mkna

	Homo sapiens activator of S phas kinase (ASK), mRNA
	Tione out to a continuous 2 (NCOA2) mPNA
	Homo sapiens nuclear receptor coactivator 3 (NCOA3), mRNA
	Homo sapiens 5T4 oncofetal trophoblast glycoprotein (5T4), mRNA
NM_002069	Homo sapiens guanine nucleotide binding protein (G protein), alpha inhibiting
	activity polypeptide 1 (GNAI1), mRNA
	Homo sapiens baculoviral IAP repeat-containing 3 (BIRC3), mRNA
NM_000391	Homo sapiens ceroid-lipofuscinosis, neuronal 2, late infantile (Jansky-
	Bielschowsky disease) (CLN2), mRNA
NM_005440	Homo sapiens GTP-binding protein Rho7 (RHO7), mRNA
	Homo sapiens heat shock 70kD protein 1B (HSPA1B), mRNA
NM_005345	Homo sapiens heat shock 70kD protein 1A (HSPA1A), mRNA
NM_003545	Homo sapiens H4 histone family, member J (H4FJ), mRNA
NM_003543	Homo sapiens H4 histone family, member H (H4FH), mRNA
	Homo sapiens H4 histone family, member G (H4FG), mRNA
NM_003540	Homo sapiens H4 histone family, member C (H4FC), mRNA
	Homo sapiens H4 histone family, member B (H4FB), mRNA
NM_003538	Homo sapiens H4 histone family, member A (H4FA), mRNA
NM_005323	Homo sapiens H1 histone family, member T (testis-specific) (H1FT), mRNA
NM_003752	Homo sapiens eukaryotic translation initiation factor 3, subunit 8 (110kD) (EIF3S8), mRNA
NM 004929	Homo sapiens calbindin 1, (28kD) (CALB1), mRNA
NM 006122	Homo sapiens mannosidase, alpha, class 2A, member 2 (MAN2A2), mRNA
NM_006301	Homo sapiens mitogen-activated protein kinase kinase kinase 12 (MAP3K12), mRNA
NM 006299	Homo sapiens zinc finger protein 193 (ZNF193), mRNA
NM 006298	Homo sapiens zinc finger protein 192 (ZNF192), mRNA
NM 006385	Homo sapiens zinc finger protein 211 (ZNF211), mRNA
NM 006296	Homo sapiens vaccinia related kinase 2 (VRK2), mRNA
NM 006295	Homo sapiens valyl-tRNA synthetase 2 (VARS2), mRNA
NM 006447	Homo sapiens ubiquitin specific protease 16 (USP16), mRNA
NM_006294	Homo sapiens ubiquinol-cytochrome c reductase binding protein (UQCRB), mRNA
NM 006293	Homo sapiens TYRO3 protein tyrosine kinase (TYRO3), mRNA
NM 006311	Homo sapiens nuclear receptor co-repressor 1 (NCOR1), mRNA
NM_006291	Homo sapiens tumor necrosis factor, alpha-induced protein 2 (TNFAIP2), mRNA
NM_006290	Homo sapiens tumor necrosis factor, alpha-induced protein 3 (TNFAIP3), mRNA
NM_006288_	Homo sapiens Thy-1 cell surface antigen (THY1), mRNA
NM_006286	Homo sapiens transcription factor Dp-2 (E2F dimerization partner 2) (TFDP2),
ND 6 006006	mRNA
NM_006284	Homo sapiens TATA box binding protein (TBP)-associated factor, RNA
) Tr (00/0/10	polymerase II, H, 30kD (TAF2H), mRNA
NM_006342	Homo sapiens transforming, acidic coiled-coil containing protein 3 (TACC3), mRNA
NM_006283	Homo sapiens transforming, acidic coiled-coil containing protein 1 (TACC1), mRNA
NM 006282	Homo sapiens serine/threonine kinase 4 (STK4), mRNA
NM_006280	Homo sapiens signal sequence receptor, delta (translocon-associated protein delta) (SSR4), mRNA
ND (00/207	Homo sapiens sushi-repeat-containing protein, X chromosome (SRPX), mRNA
NM 006307	Homo sapiens serine palmitoyltransferase, long chain base subunit 1 (SPTLC1),

	7374
	mRNA
NM_006450	Homo sapiens splicing factor (45kD) (SPF45), mRNA
NM_006422	Homo sapiens A kinase (PRKA) anchor protein 3 (AKAP3), mRNA
NM_006446	Homo sapiens solute carrier family 21 (organic anion transporter), member 6 (SLC21A6), mRNA
NM_006278	Homo sapiens sialyltransferase 4C (beta-galactosidase alpha-2,3-
1414_000270	sialytransferase) (SIAT4C), mRNA
NM_006378	Homo sapiens sema domain, immunoglobulin domain (Ig), transmembrane
111_000570	domain (TM) and short cytoplasmic domain, (semaphorin) 4D (SEMA4D), mRNA
NM_006379	Homo sapiens sema domain, immunoglobulin domain (Ig), short basic domain,
	secreted, (semaphorin) 3C (SEMA3C), mRNA
NM_006274	Homo sapiens small inducible cytokine subfamily A (Cys-Cys), member 19 (SCYA19), mRNA
NM 006453	Homo sapiens transducin (beta)-like 3 (TBL3), mRNA
NM 006270	Homo sapiens related RAS viral (r-ras) oncogene homolog (RRAS), mRNA
NM 006269	Homo sapiens retinitis pigmentosa 1 (autosomal dominant) (RP1), mRNA
NM 006355	Homo sapiens ring finger protein 15 (RNF15), mRNA
NM 006315	Homo sapiens ring finger protein 3 (RNF3), mRNA
NM 006394	Homo sapiens regulated in glioma (RIG), mRNA
NM 006263	Homo sapiens proteasome (prosome, macropain) activator subunit 1 (PA28
NIVI_UU0203	alpha) (PSME1), mRNA
NM 006262	Homo sapiens peripherin (PRPH), mRNA
NM_006261	Homo sapiens prophet of Pit1, paired-like homeodomain transcription factor (PROP1), mRNA
NM 006260	Homo sapiens protein-kinase, interferon-inducible double stranded RNA
111.2_000200	dependent inhibitor (PRKRI), mRNA
NM 006259	Homo sapiens protein kinase, cGMP-dependent, type II (PRKG2), mRNA
NM 006257	Homo sapiens protein kinase C, theta (PRKCQ), mRNA
NM 006255	Homo sapiens protein kinase C, eta (PRKCH), mRNA
NM_006253	Homo sapiens protein kinase, AMP-activated, beta 1 non-catalytic subunit (PRKAB1), mRNA
NM_006252	Homo sapiens protein kinase, AMP-activated, alpha 2 catalytic subunit
	(PRKAA2), mRNA
NM_006251	Homo sapiens protein kinase, AMP-activated, alpha 1 catalytic subunit (PRKAA1), mRNA
NM 006247	Homo sapiens protein phosphatase 5, catalytic subunit (PPP5C), mRNA
NM_006246	Homo sapiens protein phosphatase 2, regulatory subunit B (B56), epsilon
NIVI_000240	isoform (PPP2R5E), mRNA
NM_006245	Homo sapiens protein phosphatase 2, regulatory subunit B (B56), delta isoform (PPP2R5D), mRNA
NM_006244	Homo sapiens protein phosphatase 2, regulatory subunit B (B56), beta isoform (PPP2R5B), mRNA
NM_006243	Homo sapiens protein phosphatase 2, regulatory subunit B (B56), alpha isoform (PPP2R5A), mRNA
NM_006241	Homo sapiens protein phosphatase 1, regulatory (inhibitor) subunit 2 (PPP1R2), mRNA
NM_006240	Homo sapiens protein phosphatase, EF hand calcium-binding domain 1 (PPEF1), mRNA
NM_006238	Homo sapiens peroxisome proliferative activated receptor, delta (PPARD), mRNA
NM 006237	Homo sapiens POU domain, class 4, transcription factor 1 (POU4F1), mRNA
14141_000237	1 Monte supretts FOO domain, office 1, amount paon theore 1 (200 x 1), made 1

	POLI I was also 2 transmission factor 2 (POLI2F2) mPNA
NM_006236	Homo sapiens POU domain, class 3, transcription factor 3 (POU3F3), mRNA
NM_006235	Homo sapiens POU domain, class 2, associating factor 1 (POU2AF1), mRNA
NM_006231	Homo sapiens polymerase (DNA directed), epsilon (POLE), mRNA
NM_006358	Homo sapiens solute carrier family 25 (mitochondrial carrier; peroxisomal
	membrane protein, 34kD), member 17 (SLC25A17), mRNA
NM_006227	Homo sapiens phospholipid transfer protein (PLTP), mRNA
NM_006226	Homo sapiens phospholipase C, epsilon (PLCE), mRNA
NM_006225_	Homo sapiens phospholipase C, delta 1 (PLCD1), mRNA
NM_006224	Homo sapiens phosphotidylinositol transfer protein (PITPN), mRNA
NM_006479	Homo sapiens RAD51-interacting protein (PIR51), mRNA
NM_006223	Homo sapiens protein (peptidyl-prolyl cis/trans isomerase) NIMA-interacting, 4 (parvulin) (PIN4), mRNA
NM 006222	Homo sapiens protein (peptidyl-prolyl cis/trans isomerase) NIMA-interacting 1-
14141_000222	like (PIN1L), mRNA
NM 006221	Homo sapiens protein (peptidyl-prolyl cis/trans isomerase) NIMA-interacting 1
	(PIN1), mRNA
NM_006218	Homo sapiens phosphoinositide-3-kinase, catalytic, alpha polypeptide
	(PIK3CA), mRNA
NM_006213	Homo sapiens phosphorylase kinase, gamma 1 (muscle) (PHKG1), mRNA
NM_006305	Homo sapiens putative human HLA class II associated protein I (PHAP1), mRNA
NM 006212	Homo sapiens 6-phosphofructo-2-kinase/fructose-2,6-biphosphatase 2
	(PFKFB2), mRNA
NM 006211	Homo sapiens proenkephalin (PENK), mRNA
NM 006209	Homo sapiens ectonucleotide pyrophosphatase/phosphodiesterase 2 (autotaxin)
- -	(ENPP2), mRNA
NM_006205	Homo sapiens phosphodiesterase 6H, cGMP-specific, cone, gamma (PDE6H), mRNA
NM_006204	Homo sapiens phosphodiesterase 6C, cGMP-specific, cone, alpha prime
l	(PDE6C), mRNA
NM_006198	Homo sapiens Purkinje cell protein 4 (PCP4), mRNA
NM_006197	Homo sapiens pericentriolar material 1 (PCM1), mRNA
NM_006195	Homo sapiens pre-B-cell leukemia transcription factor 3 (PBX3), mRNA
NM_006193	Homo sapiens paired box gene 4 (PAX4), mRNA
NM_006191	Homo sapiens proliferation-associated 2G4, 38kD (PA2G4), mRNA
NM_006189	Homo sapiens olfactory marker protein (OMP), mRNA
NM_006186	Homo sapiens nuclear receptor subfamily 4, group A, member 2 (NR4A2), mRNA
NM 006185	Homo sapiens nuclear mitotic apparatus protein 1 (NUMA1), mRNA
NM 006184	Homo sapiens nucleobindin 1 (NUCB1), mRNA
NM 006182	Homo sapiens discoidin domain receptor family, member 2 (DDR2), mRNA
NM 006180	Homo sapiens neurotrophic tyrosine kinase, receptor, type 2 (NTRK2), mRNA
NM 006372	Homo sapiens NS1-associated protein 1 (NSAP1), mRNA
NM 006177	Homo sapiens neural retina leucine zipper (NRL), mRNA
NM 006176	Homo sapiens neurogranin (protein kinase C substrate, RC3) (NRGN), mRNA
NM 006174	Homo sapiens neuropeptide Y receptor Y5 (NPY5R), mRNA
NM 006170	Homo sapiens nucleolar protein 1 (120kD) (NOL1), mRNA
NM 006169	Homo sapiens nicotinamide N-methyltransferase (NNMT), mRNA
NM 006165	Homo sapiens nuclear factor related to kappa B binding protein (NFRKB),
11112_000103	mRNA
NM 006164	Homo sapiens nuclear factor (erythroid-derived 2)-like 2 (NFE2L2), mRNA
NM 006163	Homo sapiens nucl ar factor (erythroid-derived 2), 45kD (NFE2), mRNA
1.1.1_000103	(2)

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NM_006160	Homo sapiens neurogenic differentiation 2 (NEUROD2), mRNA
NM_006158	Homo sapiens neurofilament, light polypeptide (68kD) (NEFL), mRNA
NM_006393	Homo sapiens nebulette (NEBL), mRNA Homo sapiens DNA-binding transcriptional activator (NCYM), mRNA
NM_006316	Homo sapiens DNA-oinding transcriptional activator (NCTVI), micros
NM_006153	Homo sapiens NCK adaptor protein 1 (NCK1), mRNA
NM_006424	Homo sapiens solute carrier family 34 (sodium phosphate), member 2 (SLC34A2), mRNA
NM 006317	Homo sapiens brain acid-soluble protein 1 (BASP1), mRNA
NM 006343	Homo sapiens c-mer proto-oncogene tyrosine kinase (MERTK), mRNA
NM 006457	Homo sapiens LIM protein (similar to rat protein kinase C-binding enigma)
14141_000457	(LIM), mRNA
NM_006148	Homo sapiens LIM and SH3 protein 1 (LASP1), mRNA
NM 006383	Homo sapiens DNA-dependent protein kinase catalytic subunit-interacting
-	protein 2 (KIP2), mRNA
NM_006459	Homo sapiens similar to Caenorhabditis elegans protein C42C1.9 (KEO4),
	mRNA
NM_006147	Homo sapiens interferon regulatory factor 6 (IRF6), mRNA
NM_006332	Homo sapiens interferon, gamma-inducible protein 30 (IFI30), mRNA
NM_006337	Homo sapiens microspherule protein 1 (MCRS1), mRNA
NM_006308	Homo sapiens heat shock 27kD protein 3 (HSPB3), mRNA
NM_006403	Homo sapiens enhancer of filamentation 1 (cas-like docking; Crk-associated substrate related) (HEF1), mRNA
NR 6 006142	Homo sapiens G protein-coupled receptor 19 (GPR19), mRNA
NM_006143 NM_006302	Homo sapiens glucosidase I (GCS1), mRNA
	Homo sapiens GAS2-related on chromosome 22 (GAR22), mRNA
NM_006478	Homo sapiens glioma amplified on chromosome 1 protein (leucine-rich)
NM_006338	(GAC1), mRNA
NM 006360	Homo sapiens dendritic cell protein (GA17), mRNA
NM 006329	Homo sapiens fibulin 5 (FBLN5), mRNA
NM 006404	Homo sapiens protein C receptor, endothelial (EPCR) (PROCR), mRNA
NM 006304	Homo sapiens Deleted in split-hand/split-foot 1 region (DSS1), mRNA
NM 001355	Homo sapiens D-dopachrome tautomerase (DDT), mRNA
NM 006139	Homo sapiens CD28 antigen (Tp44) (CD28), mRNA
NM 006371	Homo sapiens cartilage associated protein (CRTAP), mRNA
NM_006136	Homo sapiens capping protein (actin filament) muscle Z-line, alpha 2
	(CAPZA2), mRNA
NM_006448	Homo sapiens trinucleotide repeat containing 1 (TNRC1), mRNA
NM_006333	Homo sapiens nuclear DNA-binding protein (C1D), mRNA
NM_006419	Homo sapiens small inducible cytokine B subfamily (Cys-X-Cys motif), member 13 (B-cell chemoattractant) (SCYB13), mRNA
NM 005453	Homo sapiens zinc finger protein 297 (ZNF297), mRNA
NM 006324	Homo sapiens craniofacial development protein 1 (CFDP1), mRNA
NM 006375	Homo sapiens cytosolic ovarian carcinoma antigen 1 (COVA1), mRNA
NM 004466	Homo sapiens glypican 5 (GPC5), mRNA
NM 004484	Homo sapiens glypican 3 (GPC3), mRNA
NM_002856	Homo sapiens poliovirus receptor-related 2 (herpesvirus entry mediator B)
	(PVRL2), mRNA
NM_001420	Homo sapiens ELAV (embryonic lethal, abnormal vision, Drosophila)-like 3 (Hu
376 001601	antigen C) (ELAVL3), mRNA
NM_001634	Homo sapiens S-adenosylmethionine decarboxylase 1 (AMD1), mRNA
NM_000483	Homo sapiens apolipoprotein C-II (APOC2), mRNA
NM_001645	Homo sapiens apolipoprotein C-I (APOC1), mRNA

NM_000482	Homo sapiens apolipoprotein A-IV (APOA4), mRNA
NM_005953	Homo sapiens metallothionein 2A (MT2A), mRNA
NM_005954	Homo sapiens metallothionein 3 (growth inhibitory factor (neurotrophic)) (MT3), mRNA
NM_006007	Homo sapiens zinc finger protein 216 (ZNF216), mRNA
NM_006006	Homo sapiens zinc finger protein 145 (Kruppel-like, expressed in promyelocytic leukemia) (ZNF145), mRNA
NM_006004	Homo sapiens ubiquinol-cytochrome c reductase hinge protein (UQCRH), mRNA
NM_006003	Homo sapiens ubiquinol-cytochrome c reductase, Rieske iron-sulfur polypeptide 1 (UQCRFS1), nuclear gene encoding mitochondrial protein, mRNA
NM_006088	Homo sapiens tubulin, beta, 2 (TUBB2), mRNA
NM_005999	Homo sapiens translin-associated factor X (TSNAX), mRNA
NM_006022	Homo sapiens transforming growth factor beta-stimulated protein TSC-22 (TSC22), mRNA
NM_005998	Homo sapiens chaperonin containing TCP1, subunit 3 (gamma) (CCT3), mRNA
NM_006073	Homo sapiens triadin (TRDN), mRNA
NM_005997	Homo sapiens transcription factor-like 1 (TCFL1), mRNA
NM_006116	Homo sapiens transforming growth factor beta-activated kinase-binding protein 1 (TAB1), mRNA
NM_005989	Homo sapiens aldo-keto reductase family 1, member D1 (delta 4-3-ketosteroid-5-beta-reductase) (AKR1D1), mRNA
NM_005988	Homo sapiens small proline-rich protein 2A (SPRR2A), mRNA
NM_005986	Homo sapiens SRY (sex determining region Y)-box 1 (SOX1), mRNA
NM_006049	Homo sapiens small nuclear RNA activating complex, polypeptide 5, 19kD (SNAPC5), mRNA
NM_006080	Homo sapiens sema domain, immunoglobulin domain (Ig), short basic domain, secreted, (semaphorin) 3A (SEMA3A), mRNA
NM_006072	Homo sapiens small inducible cytokine subfamily A (Cys-Cys), member 26 (SCYA26), mRNA
NM_005981	Homo sapiens sarcoma amplified sequence (SAS), mRNA
NM_006054	Homo sapiens reticulon 3 (RTN3), mRNA
NM_005977	Homo sapiens ring finger protein (C3H2C3 type) 6 (RNF6), mRNA
NM_005975	Homo sapiens PTK6 protein tyrosine kinase 6 (PTK6), mRNA
NM_005972	Homo sapiens pancreatic polypeptide receptor 1 (PPYR1), mRNA
NM_006112	Homo sapiens peptidylprolyl isomerase E (cyclophilin E) (PPIE), mRNA
NM_006107	Homo sapiens acid-inducible phosphoprotein (OA48-18), mRNA
NM_006067	described with medicine phosphophotom (Office 10); mid (1)
	Homo sapiens neighbor of COX4 (NOC4), mRNA
NM_005969	
NM_005969 NM_006058	Homo sapiens neighbor of COX4 (NOC4), mRNA
	Homo sapiens neighbor of COX4 (NOC4), mRNA Homo sapiens nucleosome assembly protein 1-like 4 (NAP1L4), mRNA
NM_006058 NM_006097 NM_005955	Homo sapiens neighbor of COX4 (NOC4), mRNA Homo sapiens nucleosome assembly protein 1-like 4 (NAP1L4), mRNA Homo sapiens Nef-associated factor 1 (NAF1), mRNA Homo sapiens myosin regulatory light chain 2, smooth muscle isoform (MYRL2), mRNA
NM_006058 NM_006097	Homo sapiens neighbor of COX4 (NOC4), mRNA Homo sapiens nucleosome assembly protein 1-like 4 (NAP1L4), mRNA Homo sapiens Nef-associated factor 1 (NAF1), mRNA Homo sapiens myosin regulatory light chain 2, smooth muscle isoform (MYRL2), mRNA Homo sapiens metal-regulatory transcription factor 1 (MTF1), mRNA Homo sapiens mitochondrial intermediate peptidase (MIPEP), nuclear gene
NM_006058 NM_006097 NM_005955	Homo sapiens neighbor of COX4 (NOC4), mRNA Homo sapiens nucleosome assembly protein 1-like 4 (NAP1L4), mRNA Homo sapiens Nef-associated factor 1 (NAF1), mRNA Homo sapiens myosin regulatory light chain 2, smooth muscle isoform (MYRL2), mRNA Homo sapiens metal-regulatory transcription factor 1 (MTF1), mRNA Homo sapiens mitochondrial intermediate peptidase (MIPEP), nuclear gene encoding mitochondrial protein, mRNA
NM 006058 NM_006097 NM 005955 NM_005932	Homo sapiens neighbor of COX4 (NOC4), mRNA Homo sapiens nucleosome assembly protein 1-like 4 (NAP1L4), mRNA Homo sapiens Nef-associated factor 1 (NAF1), mRNA Homo sapiens myosin regulatory light chain 2, smooth muscle isoform (MYRL2), mRNA Homo sapiens metal-regulatory transcription factor 1 (MTF1), mRNA Homo sapiens mitochondrial intermediate peptidase (MIPEP), nuclear gene encoding mitochondrial protein, mRNA Homo sapiens MHC class I polypeptide-related sequence B (MICB), mRNA
NM_006058 NM_006097 NM_005955 NM_005932 NM_005931	Homo sapiens neighbor of COX4 (NOC4), mRNA Homo sapiens nucleosome assembly protein 1-like 4 (NAP1L4), mRNA Homo sapiens Nef-associated factor 1 (NAF1), mRNA Homo sapiens myosin regulatory light chain 2, smooth muscle isoform (MYRL2), mRNA Homo sapiens metal-regulatory transcription factor 1 (MTF1), mRNA Homo sapiens mitochondrial intermediate peptidase (MIPEP), nuclear gene encoding mitochondrial protein, mRNA Homo sapiens MHC class I polypeptide-related sequence B (MICB), mRNA Homo sapiens MHC binding factor, beta (MHCBFB), mRNA Homo sapiens meningioma expressed antigen 6 (coiled-coil proline-rich)
NM 006058 NM_006097 NM 005955 NM_005932 NM 005931 NM 006081 NM_005930	Homo sapiens neighbor of COX4 (NOC4), mRNA Homo sapiens nucleosome assembly protein 1-like 4 (NAP1L4), mRNA Homo sapiens Nef-associated factor 1 (NAF1), mRNA Homo sapiens myosin regulatory light chain 2, smooth muscle isoform (MYRL2), mRNA Homo sapiens metal-regulatory transcription factor 1 (MTF1), mRNA Homo sapiens mitochondrial intermediate peptidase (MIPEP), nuclear gene encoding mitochondrial protein, mRNA Homo sapiens MHC class I polypeptide-related sequence B (MICB), mRNA Homo sapiens MHC binding factor, beta (MHCBFB), mRNA Homo sapiens meningioma expressed antigen 6 (coiled-coil proline-rich) (MGEA6), mRNA
NM_006058 NM_006097 NM_005955 NM_005932 NM_005931 NM_006081	Homo sapiens neighbor of COX4 (NOC4), mRNA Homo sapiens nucleosome assembly protein 1-like 4 (NAP1L4), mRNA Homo sapiens Nef-associated factor 1 (NAF1), mRNA Homo sapiens myosin regulatory light chain 2, smooth muscle isoform (MYRL2), mRNA Homo sapiens metal-regulatory transcription factor 1 (MTF1), mRNA Homo sapiens mitochondrial intermediate peptidase (MIPEP), nuclear gene encoding mitochondrial protein, mRNA Homo sapiens MHC class I polypeptide-related sequence B (MICB), mRNA Homo sapiens MHC binding factor, beta (MHCBFB), mRNA Homo sapiens meningioma expressed antigen 6 (coiled-coil proline-rich)

NM_005924	Homo sapiens mesenchyme homeo box 2 (growth arrest-specific homeo box) (MEOX2), mRNA
ND (005020	Homo sapiens MADS box transcription enhancer factor 2, polypeptide D
NM_005920	(myocyte enhancer factor 2D) (MEF2D), mRNA
NM 005919	Homo sapiens MADS box transcription enhancer factor 2, polypeptide B
	(myocyte enhancer factor 2B) (MEF2B), mRNA
NM_005918	Homo sapiens malate dehydrogenase 2, NAD (mitochondrial) (MDH2), nuclear
_	gene encoding mitochondrial protein, mRNA
NM 005917	Homo sapiens malate dehydrogenase 1, NAD (soluble) (MDH1), mRNA
NM 005913	Homo sapiens melanocortin 5 receptor (MC5R), mRNA
NM_005912	Homo sapiens melanocortin 4 receptor (MC4R), mRNA
NM 005911	Homo sapiens methionine adenosyltransferase II, alpha (MAT2A), mRNA
NM 005908	Homo sapiens mannosidase, beta A, lysosomal (MANBA), mRNA
NM 005907	Homo sapiens mannosidase, alpha, class 1A, member 1 (MAN1A1), mRNA
NM_005898	Homo sapiens membrane component, chromosome 11, surface marker 1
_	(M11S1), mRNA
NM 006060	Homo sapiens zinc finger protein, subfamily 1A, 1 (Ikaros) (ZNFN1A1), mRNA
NM 006059	Homo sapiens laminin, gamma 3 (LAMC3), mRNA
NM 006038	Homo sapiens spermatogenesis associated PD1 (KIAA0757), mRNA
NM_006084	Homo sapiens interferon-stimulated transcription factor 3, gamma (48kD)
_	(ISGF3G), mRNA
NM_005897	Homo sapiens intracisternal A particle-promoted polypeptide (IPP), mRNA
NM 005896	Homo sapiens isocitrate dehydrogenase 1 (NADP+), soluble (IDH1), mRNA
NM 006028	Homo sapiens 5-hydroxytryptamine (serotonin) receptor 3B (HTR3B), mRNA
NM 006120	Homo sapiens major histocompatibility complex, class II, DM alpha (HLA-
_	DMA), mRNA
NM_006026	Homo sapiens H1 histone family, member X (H1FX), mRNA
NM_006051	Homo sapiens FE65-LIKE 2 (FE65L2), mRNA
NM_006079	Homo sapiens Cbp/p300-interacting transactivator, with Glu/Asp-rich carboxy-terminal domain, 2 (CITED2), mRNA
NM_005894	Homo sapiens CD5 antigen-like (scavenger receptor cysteine rich family)
11112_00505	(CD5L), mRNA
NM_006016	Homo sapiens CD164 antigen, sialomucin (CD164), mRNA
NM_006078	Homo sapiens calcium channel, voltage-dependent, gamma subunit 2
	(CACNG2), mRNA
NM_006030	Homo sapiens calcium channel, voltage-dependent, alpha 2/delta subunit 2
	(CACNA2D2), mRNA
NM 006085	Homo sapiens 3'(2'), 5'-bisphosphate nucleotidase 1 (BPNT1), mRNA
NM 006015	Homo sapiens SWI/SNF related, matrix associated, actin dependent regulator of
_	chromatin, subfamily f, member 1 (SMARCF1), mRNA
NM_006066	Homo sapiens aldo-keto reductase family 1, member A1 (aldehyde reductase)
_	(AKR1A1), mRNA
NM_005891	Homo sapiens acetyl-Coenzyme A acetyltransferase 2 (acetoacetyl Coenzyme A
	thiolase) (ACAT2), mRNA
NM_006020	Homo sapiens alkylation repair, alkB homolog (ABH), mRNA
NM_004056	Homo sapiens carbonic anhydrase VIII (CA8), mRNA
NM_005664	Homo sapiens makorin, ring finger protein, 3 (MKRN3), mRNA
NM_005662	Homo sapiens voltage-dependent anion channel 3 (VDAC3), mRNA
NM_005836	Homo sapiens translational inhibitor protein p14.5 (UK114), mRNA
NM_005660	Homo sapiens solute carrier family 35 (UDP-galactose transporter), member 2 (SLC35A2), mRNA
NM 005659	Homo sapiens ubiquitin fusion degradation 1-like (UFD1L), mRNA

NM_005706	Homo sapiens tumor suppressing subtransferable candidate 4 (TSSC4), mRNA
NM 005723	Homo sapiens tetraspan 5 (TSPAN-5), mRNA
NM 005727	Homo sapiens tetraspan 1 (TSPAN-1), mRNA
NM 005658	Homo sapiens TNF receptor-associated factor 1 (TRAF1), mRNA
NM 005802	Homo sapiens tumor protein p53-binding protein (TP53BPL), mRNA
NM 005749	Homo sapiens transducer of ERBB2, 1 (TOB1), mRNA
NM 005655	Homo sapiens TGFB inducible early growth response (TIEG), mRNA
NM 005653	Homo sapiens transcription factor CP2 (TFCP2), mRNA
NM 005654	Homo sapiens nuclear receptor subfamily 2, group F, member 1 (NR2F1),
1.1	mRNA
NM_005652	Homo sapiens telomeric repeat binding factor 2 (TERF2), mRNA
NM 005885	Homo sapiens similar to S. cerevisiae SSM4 (TEB4), mRNA
NM 005651	Homo sapiens tryptophan 2,3-dioxygenase (TDO2), mRNA
NM 005649	Homo sapiens transcription factor 17 (TCF17), mRNA
NM 005647	Homo sapiens transducin (beta)-like 1 (TBL1), mRNA
NM_005645	Homo sapiens TATA box binding protein (TBP)-associated factor, RNA
	polymerase II, K, 18kD (TAF2K), mRNA
NM 005643	Homo sapiens TATA box binding protein (TBP)-associated factor, RNA
	polymerase II, I, 28kD (TAF2I), mRNA
NM 005641	Homo sapiens TATA box binding protein (TBP)-associated factor, RNA
	polymerase II, E, 70/85kD (TAF2E), mRNA
NM 005679	Homo sapiens TATA box binding protein (TBP)-associated factor, RNA
-	polymerase I, C, 110kD (TAF1C), mRNA
NM 005681	Homo sapiens TATA box binding protein (TBP)-associated factor, RNA
-	polymerase I, A, 48kD (TAF1A), mRNA
NM 005639	Homo sapiens synaptotagmin 1 (SYT1), mRNA
NM 005638	Homo sapiens synaptobrevin-like 1 (SYBL1), mRNA
NM 005635	Homo sapiens synovial sarcoma, X breakpoint 1 (SSX1), mRNA
NM_005871	Homo sapiens splicing factor 30, survival of motor neuron-related (SPF30),
	mRNA
NM_005634	Homo sapiens SRY (sex determining region Y)-box 3 (SOX3), mRNA
NM_005686	Homo sapiens SRY (sex determining region Y)-box 13 (SOX13), mRNA
NM_005629	Homo sapiens solute carrier family 6 (neurotransmitter transporter, creatine),
	member 8 (SLC6A8), mRNA
NM_005630	Homo sapiens solute carrier family 21 (prostaglandin transporter), member 2
	(SLC21A2), mRNA
NM_005628	Homo sapiens solute carrier family 1 (neutral amino acid transporter), member 5
	(SLC1A5), mRNA
NM_005627	Homo sapiens serum/glucocorticoid regulated kinase (SGK), mRNA
NM_005877	Homo sapiens splicing factor 3a, subunit 1, 120kD (SF3A1), mRNA
NM: 005625	Homo sapiens syndecan binding protein (syntenin) (SDCBP), mRNA
NM_005623	Homo sapiens small inducible cytokine subfamily A (Cys-Cys), member 8
	(monocyte chemotactic protein 2) (SCYA8), mRNA
NM_005624	Homo sapiens small inducible cytokine subfamily A (Cys-Cys), member 25
275 005050	(SCYA25), mRNA
NM_005850	Homo sapiens splicing factor 3b, subunit 4, 49kD (SF3B4), mRNA
NM_005772	Homo sapiens RNA cyclase homolog (RNAC), mRNA
NM_005614	Homo sapiens Ras homolog enriched in brain 2 (RHEB2), mRNA
NM_005777	Homo sapiens RNA binding motif protein 6 (RBM6), mRNA
NM_005778	Homo sapiens RNA binding motif protein 5 (RBM5), mRNA
NM_005611	Homo sapiens retinoblastoma-like 2 (p130) (RBL2), mRNA
NM_005704	Homo sapiens protein tyrosine phosphatase, receptor type, U (PTPRU), mRNA

NM 005607	Homo sapiens PTK2 protein tyrosine kinase 2 (PTK2), mRNA
NM_005789	Homo sapiens proteasome (prosome, macropain) activator subunit 3 (PA28
212.2_	gamma; Ki) (PSME3), mRNA
NM 005672	Hom sapiens prostate stem cell antigen (PSCA), mRNA
NM_005865	Homo sapiens protease, serine, 16 (thymus) (PRSS16), mRNA
NM 005729	Homo sapiens peptidylprolyl isomerase F (cyclophilin F) (PPIF), mRNA
NM_005604	Homo sapiens POU domain, class 3, transcription factor 2 (POU3F2), mRNA
NM 005709	Homo sapiens PDZ-73 protein (PDZ-73/NY-CO-38), mRNA
	Homo sapiens purinergic receptor (family A group 5) (P2Y5), mRNA
NM_005767	Homo sapiens solute carrier family 17 (sodium phosphate), member 2
NM_005835	(SLC17A2), mRNA
NM_005793	Homo sapiens nucleoside diphosphate kinase type 6 (inhibitor of p53-induced
	apoptosis-alpha) (NM23-H6), mRNA
NM_005600	Homo sapiens nitrilase 1 (NIT1), mRNA
NM_005599	Homo sapiens nescient helix loop helix 2 (NHLH2), mRNA
NM_005598	Homo sapiens nescient helix loop helix 1 (NHLH1), mRNA
NM_005596	Homo sapiens nuclear factor I/B (NFIB), mRNA
NM_005665	Homo sapiens ecotropic viral integration site 5 (EVI5), mRNA
NM_005594	Homo sapiens nascent-polypeptide-associated complex alpha polypeptide (NACA), mRNA
NM 005593	Homo sapiens myogenic factor 5 (MYF5), mRNA
NM 005592	Homo sapiens muscle, skeletal, receptor tyrosine kinase (MUSK), mRNA
NM_005845	Homo sapiens ATP-binding cassette, sub-family C (CFTR/MRP), member 4
	(ABCC4), mRNA
NM_005874	Homo sapiens leukocyte immunoglobulin-like receptor, subfamily B (with TM and ITIM domains), member 2 (LILRB2), mRNA
NM_005588	Homo sapiens meprin A, alpha (PABA peptide hydrolase) (MEP1A), mRNA
NM_005587	Homo sapiens MADS box transcription enhancer factor 2, polypeptide A (myocyte enhancer factor 2A) (MEF2A), mRNA
NM_005810	Homo sapiens killer cell lectin-like receptor subfamily G, member 1 (KLRG1), mRNA
NM 005581	Homo sapiens Lutheran blood group (Auberger b antigen included) (LU), mRNA
NM_005578	Homo sapiens LIM domain-containing preferred translocation partner in lipoma (LPP), mRNA
NM 005577	Homo sapiens lipoprotein, Lp(a) (LPA), mRNA
NM 005576	Homo sapiens lysyl oxidase-like 1 (LOXL1), mRNA
NM 005573	Homo sapiens lamin B1 (LMNB1), mRNA
NM_005572	Homo sapiens lamin A/C (LMNA), mRNA
NM 005568	Homo sapiens LIM homeobox protein 1 (LHX1), mRNA
NM 005780	Homo sapiens lipoma HMGIC fusion partner (LHFP), mRNA
NM 005566	Homo sapiens lactate dehydrogenase A (LDHA), mRNA
	Homo sapiens lipocalin 2 (oncogene 24p3) (LCN2), mRNA
NM_005564	Homo sapiens ladinin 1 (LAD1), mRNA
NM 005558	Homo sapiens keratin 7 (KRT7), mRNA
NM_005556	Homo sapiens keratin 16 (focal non-epidermolytic palmoplantar keratoderma)
NM_005557	(KRT16), mRNA
NM 005553	Homo sapiens keratin, cuticle, ultrahigh sulphur 1 (KRN1), mRNA
NM 005552	Homo sapiens kinesin 2 (60-70kD) (KNS2), mRNA
NM 005551	Homo sapiens kallikrein 2, prostatic (KLK2), mRNA
NM 005550	Homo sapiens kinesin family member C3 (KIFC3), mRNA
	Homo sapiens potassium large conductance calcium-activated channel,
NM_005832	subfamily M, beta member 2 (KCNMB2), mRNA

NM_005549	Homo sapiens potassium voltage-gated channel, shaker-related subfamily, member 10 (KCNA10), mRNA
NM_005548	Homo sapiens lysyl-tRNA synthetase (KARS), mRNA
NM 005547	Homo sapiens involucrin (IVL), mRNA
NM 005545	Homo sapiens immunoglobulin superfamily containing leucine-rich repeat
1414_005545	(ISLR), mRNA
NM_005853	Homo sapiens iroquois-class homeodomain protein (IRX-2A), mRNA
NM_005544	Homo sapiens insulin receptor substrate 1 (IRS1), mRNA
NM_005543	Homo sapiens insulin-like 3 (Leydig cell) (INSL3), mRNA
NM_005542	Homo sapiens insulin induced gene 1 (INSIG1), mRNA
NM_005541	Homo sapiens inositol polyphosphate-5-phosphatase, 145kD (INPP5D), mRNA
NM_005539	Homo sapiens inositol polyphosphate-5-phosphatase, 40kD (INPP5A), mRNA
NM_005537	Homo sapiens inhibitor of growth 1 family, member 1 (ING1), mRNA
NM_005535	Homo sapiens interleukin 12 receptor, beta 1 (IL12RB1), mRNA
NM 005532	Homo sapiens interferon, alpha-inducible protein 27 (IFI27), mRNA
NM_005531	Homo sapiens interferon, gamma-inducible protein 16 (IFI16), mRNA
NM_005530	Homo sapiens isocitrate dehydrogenase 3 (NAD+) alpha (IDH3A), mRNA
NM_005808	Homo sapiens HYA22 protein (HYA22), mRNA
NM_005528	Homo sapiens heat shock 40kD protein 2 (HSPF2), mRNA
NM_005526	Homo sapiens heat shock transcription factor 1 (HSF1), mRNA
NM_005525	Homo sapiens hydroxysteroid (11-beta) dehydrogenase 1 (HSD11B1), mRNA
NM_005522	Homo sapiens homeo box A1 (HOXA1), mRNA
NM_005521	Homo sapiens homeo box 11 (T-cell lymphoma 3-associated breakpoint)
	(HOX11), mRNA
NM_005518	Homo sapiens 3-hydroxy-3-methylglutaryl-Coenzyme A synthase 2
	(mitochondrial) (HMGCS2), mRNA
NM_005515	Homo sapiens homeo box HB9 (HLXB9), mRNA
NM_005516	Homo sapiens major histocompatibility complex, class I, E (HLA-E), mRNA
NM_005712	Homo sapiens HERV-H LTR-associating 1 (HHLA1), mRNA
NM_005844	Homo sapiens PERB11 family member in MHC class I region (HCGIX), mRNA
NM_005513	Homo sapiens general transcription factor IIE, polypeptide 1 (alpha subunit, 56kD) (GTF2E1), mRNA
NM 005683	Homo sapiens G protein-coupled receptor 55 (GPR55), mRNA
NM 005684	Homo sapiens G protein-coupled receptor 52 (GPR52), mRNA
NM 005512	Homo sapiens glycoprotein A repetitions predominant (GARP), mRNA
NM_005851	Homo sapiens tumor suppressor deleted in oral cancer-related 1 (DOC-1R), mRNA
NM_005740	Homo sapiens dynein, axonemal, light polypeptide 4 (DNAL4), mRNA
NM 005872	Homo sapiens breast carcinoma amplified sequence 2 (BCAS2), mRNA
NM 005671	Homo sapiens reproduction 8 (D8S2298E), mRNA
NM_005800	Homo sapiens highly charged protein (D13S106E), mRNA
NM 005752	Homo sapiens C-type (calcium dependent, carbohydrate-recognition domain)
1111_003/32	lectin, superfamily member 1 (cartilage-derived) (CLECSF1), mRNA
NM 005507	Homo sapiens cofilin 1 (non-muscle) (CFL1), mRNA
NM_005825	Homo sapiens RAS guanyl releasing protein 2 (calcium and DAG-regulated)
1111_000025	(RASGRP2), mRNA
NM 005773	Homo sapiens zinc finger protein 256 (ZNF256), mRNA
NM 005774	Homo sapiens zinc finger protein 255 (ZNF255), mRNA
NM 005504	Homo sapiens branched chain aminotransferase 1, cytosolic (BCAT1), mRNA
NM 005738	Homo sapiens ADP-ribosylation factor-like 4 (ARL4), mRNA
NM_005731	Homo sapiens actin related protein 2/3 complex, subunit 2 (34 kD) (ARPC2),
	mRNA

NM_005719	Homo sapiens actin related protein 2/3 complex, subunit 3 (21 kD) (ARPC3), mRNA
NM 005883	Homo sapiens adenomatous polyposis coli like (APCL), mRNA
NM 005858	Homo sapiens A kinase (PRKA) anchor protein 8 (AKAP8), mRNA
NM 002023	Homo sapiens fibromodulin (FMOD), mRNA
NM_000108	Homo sapiens dihydrolipoamide dehydrogenase (E3 component of pyruvate
11111_000100	dehydrogenase complex, 2-oxo-glutarate complex, branched chain keto acid
	dehydrogenase complex) (DLD), mRNA
NM 001621	Homo sapiens aryl hydrocarbon receptor (AHR), mRNA
NM 001101	Homo sapiens actin, beta (ACTB), mRNA
NM_001100	Homo sapiens actin, alpha 1, skeletal muscle (ACTA1), mRNA
NM 000054	Homo sapiens arginine vasopressin receptor 2 (nephrogenic diabetes insipidus)
11112_00000	(AVPR2), mRNA
NM 005455	Homo sapiens zinc finger protein 265 (ZNF265), mRNA
NM 005433	Homo sapiens v-yes-1 Yamaguchi sarcoma viral oncogene homolog 1 (YES1),
	mRNA
NM 005429	Homo sapiens vascular endothelial growth factor C (VEGFC), mRNA
NM 005499	Homo sapiens SUMO-1 activating enzyme subunit 2 (UBA2), mRNA
NM 005427	Homo sapiens tumor protein p73 (TP73), mRNA
NM 005425	Homo sapiens transition protein 2 (during histone to protamine replacement)
	(TNP2), mRNA
NM 005424	Homo sapiens tyrosine kinase with immunoglobulin and epidermal growth factor
	homology domains (TIE), mRNA
NM 005423	Homo sapiens trefoil factor 2 (spasmolytic protein 1) (TFF2), mRNA
NM 005422	Homo sapiens tectorin alpha (TECTA), mRNA
NM 005421	Homo sapiens T-cell acute lymphocytic leukemia 2 (TAL2), mRNA
NM_005420	Homo sapiens sulfotransferase, estrogen-preferring (STE), mRNA
NM 005418	Homo sapiens suppression of tumorigenicity 5 (ST5), mRNA
NM 005470	Homo sapiens spectrin SH3 domain binding protein 1 (SSH3BP1), mRNA
NM 005416	Homo sapiens small proline-rich protein 3 (SPRR3), mRNA
NM 005460	Homo sapiens synuclein, alpha interacting protein (synphilin) (SNCAIP), mRNA
NM_005412	Homo sapiens serine hydroxymethyltransferase 2 (mitochondrial) (SHMT2), mRNA
NM_005408	Homo sapiens small inducible cytokine subfamily A (Cys-Cys), member 13 (SCYA13), mRNA
NM_005402	Homo sapiens v-ral simian leukemia viral oncogene homolog A (ras related) (RALA), mRNA
NM 005397	Homo sapiens podocalyxin-like (PODXL), mRNA
NM 005395	Homo sapiens postmeiotic segregation increased 2-like 9 (PMS2L9), mRNA
NM 005394	Homo sapiens postmeiotic segregation increased 2-like 8 (PMS2L8), mRNA
NM 005390	Homo sapiens pyruvate dehydrogenase (lipoamide) alpha 2 (PDHA2), mRNA
NM 005389	Homo sapiens protein-L-isoaspartate (D-aspartate) O-methyltransferase
	(PCMT1), mRNA
NM_005450	Homo sapiens noggin (NOG), mRNA
NM_005386	Homo sapiens neuronatin (NNAT), mRNA
NM 005384	Homo sapiens nuclear factor, interleukin 3 regulated (NFIL3), mRNA
NM 005383	Homo sapiens sialidase 2 (cytosolic sialidase) (NEU2), mRNA
NM 005382	Homo sapiens neurofilament 3 (150kD medium) (NEF3), mRNA
NM 005381	Homo sapiens nucleolin (NCL), mRNA
NM 005380	Homo sapiens neuroblastoma, suppression of tumorigenicity 1 (NBL1), mRNA
NM_005468	Homo sapiens N-acetylated alpha-linked acidic dipeptidase-like; ILEAL
	DIPEPTIDYLPEPTIDASE (NAALADASEL), mRNA

NM_005374	Homo sapiens membrane protein, palmitoylated 2 (MAGUK p55 subfamily member 2) (MPP2), mRNA
NM 005373	Homo sapiens myeloproliferative leukemia virus oncogene (MPL), mRNA
NM_005372	Homo sapiens v-mos Moloney murine sarcoma viral oncogene homolog (MOS),
14141_005572	mRNA
NM 005439	Homo sapiens myeloid leukemia factor 2 (MLF2), mRNA
NM 005369	Homo sapiens MCF.2 cell line derived transforming sequence (MCF2), mRNA
NM 005368	Homo sapiens myoglobin (MB), mRNA
NM 005363	Homo sapiens melanoma antigen, family A, 6 (MAGEA6), mRNA
NM 005362	Homo sapiens melanoma antigen, family A, 3 (MAGEA3), mRNA
NM 005361	Homo sapiens melanoma antigen, family A, 2 (MAGEA2), mRNA
NM 005475	Homo sapiens lymphocyte adaptor protein (LNK), mRNA
NM 005357	Homo sapiens lipase, hormone-sensitive (LIPE), mRNA
NM 005356	Homo sapiens lymphocyte-specific protein tyrosine kinase (LCK), mRNA
NM_005472	Homo sapiens potassium voltage-gated channel, Isk-related family, member 3
1441_005.72	(KCNE3), mRNA
NM 005495	Homo sapiens solute carrier family 17 (sodium phosphate), member 4
	(SLC17A4), mRNA
NM 005456	Homo sapiens mitogen-activated protein kinase 8 interacting protein 1
11212_000 100	(MAPK 8IP1), mRNA
NM 005343	Homo sapiens v-Ha-ras Harvey rat sarcoma viral oncogene homolog (HRAS),
	mRNA ·
NM_005342	Homo sapiens high-mobility group (nonhistone chromosomal) protein 4
- · · -	(HMG4), mRNA
NM 005341	Homo sapiens GLI-Kruppel family member HKR3 (HKR3), mRNA
NM 005337	Homo sapiens hematopoietic protein 1 (HEM1), mRNA
NM 005477	Homo sapiens hyperpolarization activated cyclic nucleotide-gated potassium
_	channel 4 (HCN4), mRNA
NM_005335	Homo sapiens hematopoietic cell-specific Lyn substrate 1 (HCLS1), mRNA
NM_005334	Homo sapiens host cell factor C1 (VP16-accessory protein) (HCFC1), mRNA
NM_005333	Homo sapiens holocytochrome c synthase (cytochrome c heme-lyase) (HCCS),
	mRNA
NM_005328	Homo sapiens hyaluronan synthase 2 (HAS2), mRNA
NM_005327	Homo sapiens L-3-hydroxyacyl-Coenzyme A dehydrogenase, short chain
	(HADHSC), mRNA
NM_005324	Homo sapiens H3 histone, family 3B (H3.3B) (H3F3B), mRNA
NM_005321	Homo sapiens H1 histone family, member 4 (H1F4), mRNA
NM_005320	Homo sapiens H1 histone family, member 3 (H1F3), mRNA
NM_005319	Homo sapiens H1 histone family, member 2 (H1F2), mRNA
NM_005325	Homo sapiens H1 histone family, member 1 (H1F1), mRNA
NM_005318	Homo sapiens H1 histone family, member 0 (H1F0), mRNA
NM_005459	Homo sapiens guanylate cyclase activator 1C (GUCA1C), mRNA
NM_005316	Homo sapiens general transcription factor IIH, polypeptide 1 (62kD subunit)
	(GTF2H1), mRNA
NM_005315	Homo sapiens goosecoid-like (GSCL), mRNA
NM_005314	Homo sapiens gastrin-releasing peptide receptor (GRPR), mRNA
NM_005313	Homo sapiens glucose regulated protein, 58kD (GRP58), mRNA
NM_005312	Homo sapiens guanine nucleotide-releasing factor 2 (specific for crk proto-
	oncogene) (GRF2), mRNA
NM_005311	Homo sapiens growth factor receptor-bound protein 10 (GRB10), mRNA
NM_005309	Homo sapiens glutamic-pyruvate transaminase (alanine aminotransferase)
	(GPT), mRNA

NM_005308	Homo sapiens G protein-coupled receptor kinase 5 (GPRK5), mRNA
NM_005286	Homo sapiens G protein-coupled receptor 8 (GPR8), mRNA
NM_005285	Homo sapiens G protein-coupled receptor 7 (GPR7), mRNA
NM_005284	Homo sapiens G protein-coupled receptor 6 (GPR6), mRNA
NM_005458	Homo sapiens G protein-coupled receptor 51 (GPR51), mRNA
NM 005282	Homo sapiens G protein-coupled receptor 4 (GPR4), mRNA
NM 005306	Homo sapiens G protein-coupled receptor 43 (GPR43), mRNA
NM 005305	Homo sapiens G protein-coupled receptor 42 (GPR42), mRNA
NM 005304	Homo sapiens G protein-coupled receptor 41 (GPR41), mRNA
NM 005303	Homo sapiens G protein-coupled receptor 40 (GPR40), mRNA
NM 005281	Homo sapiens G protein-coupled receptor 3 (GPR3), mRNA
NM 005302	Homo sapiens G protein-coupled receptor 37 (endothelin receptor type B-like)
	(GPR37), mRNA
NM 005301	Homo sapiens G protein-coupled receptor 35 (GPR35), mRNA
NM 005300	Homo sapiens G protein-coupled receptor 34 (GPR34), mRNA
NM 005299	Homo sapiens G protein-coupled receptor 31 (GPR31), mRNA
NM 005298	Homo sapiens G protein-coupled receptor 25 (GPR25), mRNA
NM 005297	Homo sapiens G protein-coupled receptor 24 (GPR24), mRNA
NM 005296	Homo sapiens G protein-coupled receptor 23 (GPR23), mRNA
NM 005295	Homo sapiens G protein-coupled receptor 22 (GPR22), mRNA
NM_005294	Homo sapiens G protein-coupled receptor 21 (GPR21), mRNA
NM 005293	Homo sapiens G protein-coupled receptor 20 (GPR20), mRNA
NM 005279	Homo sapiens G protein-coupled receptor 1 (GPR1), mRNA
NM 005291	Homo sapiens G protein-coupled receptor 17 (GPR17), mRNA
NM 005290	Homo sapiens G protein-coupled receptor 15 (GPR15), mRNA
NM 005288	Homo sapiens G protein-coupled receptor 12 (GPR12), mRNA
NM 005276	Homo sapiens glycerol-3-phosphate dehydrogenase 1 (soluble) (GPD1), mRNA
NM 005275	Homo sapiens guanine nucleotide binding protein-like 1 (GNL1), mRNA
NM 005274	Homo sapiens guanine nucleotide binding protein (G protein), gamma 5
14141_005274	(GNG5), mRNA
NM 005273	Homo sapiens guanine nucleotide binding protein (G protein), beta polypeptide 2
1411_003273	(GNB2), mRNA
NM 005271	Homo sapiens glutamate dehydrogenase 1 (GLUD1), mRNA
NM_005269	Homo sapiens glioma-associated oncogene homolog (zinc finger protein) (GLI),
11112_000203	mRNA
NM 005264	Homo sapiens GDNF family receptor alpha 1 (GFRA1), mRNA
NM 005263	Homo sapiens growth factor independent 1 (GFI1), mRNA
NM_005256	Homo sapiens growth arrest-specific 2 (GAS2), mRNA
NM_005255	Homo sapiens cyclin G associated kinase (GAK), mRNA
NM 005253	Homo sapiens FOS-like antigen 2 (FOSL2), mRNA
NM_005249	Homo sapiens forkhead box G1B (FOXG1B), mRNA
NM_005251	Homo sapiens forkhead box C2 (MFH-1, mesenchyme forkhead 1) (FOXC2),
1111_005251	mRNA
NM_005248	Homo sapiens Gardner-Rasheed feline sarcoma viral (v-fgr) oncogene homolog
1111_005270	(FGR), mRNA
NM 005246	Homo sapiens fer (fps/fes related) tyrosine kinase (phosphoprotein NCP94)
11111_005270	(FER), mRNA
NM_005234	Homo sapiens nuclear receptor subfamily 2, group F, member 6 (NR2F6),
11171_005254	mRNA
NM 005233	Homo sapiens EphA3 (EPHA3), mRNA
NM_005233	Homo sapiens ems1 sequence (mammary tumor and squamous cell carcinoma-
1111_005251	associated (p80/85 src substrate) (EMS1), mRNA
L	associated (box or sto succeed) (Dirior), mader

apiens ephrin-A4 (EFNA4), mRNA
apiens d oxyribonuclease I (DNASE1), mRNA
apiens distal-less homeo box 6 (DLX6), mRNA
apiens distal-less homeo box 3 (DLX3), mRNA
sapiens dolichyl-diphosphooligosaccharide-protein glycosyltransferase
T), mRNA
sapiens deleted in colorectal carcinoma (DCC), mRNA
sapiens DNA segment, single copy, probe pH4 (transforming sequence,
-1, (D10S170), mRNA
sapiens cytotoxic T-lymphocyte-associated protein 4 (CTLA4), mRNA
sapiens cystatin A (stefin A) (CSTA), mRNA
sapiens cystatin 8 (cystatin-related epididymal specific) (CST8), mRNA
sapiens casein, kappa (CSN10), mRNA
sapiens colony stimulating factor 1 receptor, formerly McDonough feline
a viral (v-fms) oncogene homolog (CSF1R), mRNA
sapiens mitogen-activated protein kinase kinase kinase 8 (MAP3K8),
sapiens cell matrix adhesion regulator (CMAR), mRNA
sapiens CCAAT/enhancer binding protein (C/EBP), delta (CEBPD),
sapiens CCAAT/enhancer binding protein (C/EBP), beta (CEBPB),
sapiens caudal type homeo box transcription factor 4 (CDX4), mRNA
sapiens CD80 antigen (CD28 antigen ligand 1, B7-1 antigen) (CD80),
sapiens Cas-Br-M (murine) ecotropic retroviral transforming sequence
mRNA
sapiens calmodulin-like 3 (CALML3), mRNA
sapiens calmodulin 3 (phosphorylase kinase, delta) (CALM3), mRNA
sapiens chromatin assembly factor 1, subunit A (p150) (CHAF1A),
sapiens chromatin assembly factor 1, subunit B (p60) (CHAF1B), mRNA
sapiens calcium channel, voltage-dependent, alpha 1F subunit
sapiens calcium channel, voltage-dependent, alpha 1F subunit NA1F), mRNA
NA1F), mRNA sapiens carbonic anhydrase VII (CA7), mRNA
NA1F), mRNA sapiens carbonic anhydrase VII (CA7), mRNA sapiens bone morphogenetic protein 15 (BMP15), mRNA
NA1F), mRNA sapiens carbonic anhydrase VII (CA7), mRNA sapiens bone morphogenetic protein 15 (BMP15), mRNA sapiens B-cell CLL/lymphoma 3 (BCL3), mRNA
NA1F), mRNA sapiens carbonic anhydrase VII (CA7), mRNA sapiens bone morphogenetic protein 15 (BMP15), mRNA sapiens B-cell CLL/lymphoma 3 (BCL3), mRNA sapiens ATPase, H+ transporting, lysosomal (vacuolar proton pump) non-
NA1F), mRNA sapiens carbonic anhydrase VII (CA7), mRNA sapiens bone morphogenetic protein 15 (BMP15), mRNA sapiens B-cell CLL/lymphoma 3 (BCL3), mRNA sapiens ATPase, H+ transporting, lysosomal (vacuolar proton pump) non- ic accessory protein 1A (110/116kD) (ATP6N1A), mRNA
NA1F), mRNA sapiens carbonic anhydrase VII (CA7), mRNA sapiens bone morphogenetic protein 15 (BMP15), mRNA sapiens B-cell CLL/lymphoma 3 (BCL3), mRNA sapiens ATPase, H+ transporting, lysosomal (vacuolar proton pump) non- ic accessory protein 1A (110/116kD) (ATP6N1A), mRNA sapiens ATP synthase, H+ transporting, mitochondrial F1 complex,
NA1F), mRNA sapiens carbonic anhydrase VII (CA7), mRNA sapiens bone morphogenetic protein 15 (BMP15), mRNA sapiens B-cell CLL/lymphoma 3 (BCL3), mRNA sapiens ATPase, H+ transporting, lysosomal (vacuolar proton pump) non- ic accessory protein 1A (110/116kD) (ATP6N1A), mRNA sapiens ATP synthase, H+ transporting, mitochondrial F1 complex, a polypeptide 1 (ATP5C1), mRNA
NA1F), mRNA sapiens carbonic anhydrase VII (CA7), mRNA sapiens bone morphogenetic protein 15 (BMP15), mRNA sapiens B-cell CLL/lymphoma 3 (BCL3), mRNA sapiens ATPase, H+ transporting, lysosomal (vacuolar proton pump) non- ic accessory protein 1A (110/116kD) (ATP6N1A), mRNA sapiens ATP synthase, H+ transporting, mitochondrial F1 complex, a polypeptide 1 (ATP5C1), mRNA sapiens ATPase, Ca++ transporting, ubiquitous (ATP2A3), mRNA
NA1F), mRNA sapiens carbonic anhydrase VII (CA7), mRNA sapiens bone morphogenetic protein 15 (BMP15), mRNA sapiens B-cell CLL/lymphoma 3 (BCL3), mRNA sapiens ATPase, H+ transporting, lysosomal (vacuolar proton pump) non- ic accessory protein 1A (110/116kD) (ATP6N1A), mRNA sapiens ATP synthase, H+ transporting, mitochondrial F1 complex, a polypeptide 1 (ATP5C1), mRNA sapiens ATPase, Ca++ transporting, ubiquitous (ATP2A3), mRNA sapiens activating transcription factor 1 (ATF1), mRNA
NA1F), mRNA sapiens carbonic anhydrase VII (CA7), mRNA sapiens bone morphogenetic protein 15 (BMP15), mRNA sapiens B-cell CLL/lymphoma 3 (BCL3), mRNA sapiens ATPase, H+ transporting, lysosomal (vacuolar proton pump) non- ic accessory protein 1A (110/116kD) (ATP6N1A), mRNA sapiens ATP synthase, H+ transporting, mitochondrial F1 complex, a polypeptide 1 (ATP5C1), mRNA sapiens ATPase, Ca++ transporting, ubiquitous (ATP2A3), mRNA sapiens activating transcription factor 1 (ATF1), mRNA sapiens ras homolog gene family, member C (ARHC), mRNA
sapiens carbonic anhydrase VII (CA7), mRNA sapiens bone morphogenetic protein 15 (BMP15), mRNA sapiens B-cell CLL/lymphoma 3 (BCL3), mRNA sapiens ATPase, H+ transporting, lysosomal (vacuolar proton pump) non- ic accessory protein 1A (110/116kD) (ATP6N1A), mRNA sapiens ATP synthase, H+ transporting, mitochondrial F1 complex, a polypeptide 1 (ATP5C1), mRNA sapiens ATPase, Ca++ transporting, ubiquitous (ATP2A3), mRNA sapiens activating transcription factor 1 (ATF1), mRNA sapiens ras homolog gene family, member C (ARHC), mRNA sapiens amyloid beta (A4) precursor-like protein 1 (APLP1), mRNA
sapiens carbonic anhydrase VII (CA7), mRNA sapiens bone morphogenetic protein 15 (BMP15), mRNA sapiens B-cell CLL/lymphoma 3 (BCL3), mRNA sapiens ATPase, H+ transporting, lysosomal (vacuolar proton pump) non- ic accessory protein 1A (110/116kD) (ATP6N1A), mRNA sapiens ATP synthase, H+ transporting, mitochondrial F1 complex, a polypeptide 1 (ATP5C1), mRNA sapiens ATPase, Ca++ transporting, ubiquitous (ATP2A3), mRNA sapiens activating transcription factor 1 (ATF1), mRNA sapiens ras homolog gene family, member C (ARHC), mRNA sapiens amyloid beta (A4) precursor-like protein 1 (APLP1), mRNA sapiens aldolase C, fructose-bisphosphate (ALDOC), mRNA
sapiens carbonic anhydrase VII (CA7), mRNA sapiens bone morphogenetic protein 15 (BMP15), mRNA sapiens B-cell CLL/lymphoma 3 (BCL3), mRNA sapiens ATPase, H+ transporting, lysosomal (vacuolar proton pump) non- ic accessory protein 1A (110/116kD) (ATP6N1A), mRNA sapiens ATP synthase, H+ transporting, mitochondrial F1 complex, a polypeptide 1 (ATP5C1), mRNA sapiens ATPase, Ca++ transporting, ubiquitous (ATP2A3), mRNA sapiens activating transcription factor 1 (ATF1), mRNA sapiens ras homolog gene family, member C (ARHC), mRNA sapiens amyloid beta (A4) precursor-like protein 1 (APLP1), mRNA
sapiens carbonic anhydrase VII (CA7), mRNA sapiens bone morphogenetic protein 15 (BMP15), mRNA sapiens B-cell CLL/lymphoma 3 (BCL3), mRNA sapiens ATPase, H+ transporting, lysosomal (vacuolar proton pump) non- ic accessory protein 1A (110/116kD) (ATP6N1A), mRNA sapiens ATP synthase, H+ transporting, mitochondrial F1 complex, a polypeptide 1 (ATP5C1), mRNA sapiens ATPase, Ca++ transporting, ubiquitous (ATP2A3), mRNA sapiens activating transcription factor 1 (ATF1), mRNA sapiens ras homolog gene family, member C (ARHC), mRNA sapiens amyloid beta (A4) precursor-like protein 1 (APLP1), mRNA sapiens aldolase C, fructose-bisphosphate (ALDOC), mRNA sapiens v-akt murine thymoma viral oncogene homolog 1 (AKT1),
sapiens carbonic anhydrase VII (CA7), mRNA sapiens bone morphogenetic protein 15 (BMP15), mRNA sapiens B-cell CLL/lymphoma 3 (BCL3), mRNA sapiens ATPase, H+ transporting, lysosomal (vacuolar proton pump) non- ic accessory protein 1A (110/116kD) (ATP6N1A), mRNA sapiens ATP synthase, H+ transporting, mitochondrial F1 complex, a polypeptide 1 (ATP5C1), mRNA sapiens ATPase, Ca++ transporting, ubiquitous (ATP2A3), mRNA sapiens activating transcription factor 1 (ATF1), mRNA sapiens ras homolog gene family, member C (ARHC), mRNA sapiens amyloid beta (A4) precursor-like protein 1 (APLP1), mRNA sapiens aldolase C, fructose-bisphosphate (ALDOC), mRNA sapiens v-akt murine thymoma viral oncogene homolog 1 (AKT1),
sapiens carbonic anhydrase VII (CA7), mRNA sapiens bone morphogenetic protein 15 (BMP15), mRNA sapiens B-cell CLL/lymphoma 3 (BCL3), mRNA sapiens ATPase, H+ transporting, lysosomal (vacuolar proton pump) non- ic accessory protein 1A (110/116kD) (ATP6N1A), mRNA sapiens ATP synthase, H+ transporting, mitochondrial F1 complex, a polypeptide 1 (ATP5C1), mRNA sapiens ATPase, Ca++ transporting, ubiquitous (ATP2A3), mRNA sapiens activating transcription factor 1 (ATF1), mRNA sapiens ras homolog gene family, member C (ARHC), mRNA sapiens amyloid beta (A4) precursor-like protein 1 (APLP1), mRNA sapiens aldolase C, fructose-bisphosphate (ALDOC), mRNA sapiens v-akt murine thymoma viral oncogene homolog 1 (AKT1), sapiens angiotensin receptor-like 1 (AGTRL1), mRNA

	1 1 1 2 1 2 1 2 1 2 1 2 1 2 1 2 1 2 1 2
NM_005121	Homo sapiens thyroid hormone receptor-associated protein, 240 kDa subunit (TRAP240), mRNA
NM 005079	Homo sapiens tumor protein D52 (TPD52), mRNA
NM 005091	Homo sapiens peptidoglycan recognition protein (PGLYRP), mRNA
NM_005092	Homo sapiens tumor necrosis factor (ligand) superfamily, member 18 (TNFSF18), mRNA
NM_005118	Homo sapiens tumor necrosis factor (ligand) superfamily, member 15 (TNFSF15), mRNA
NM 005147	Homo sapiens tumorous imaginal discs (Drosophila) homolog (TID1), mRNA
NM 005076	Homo sapiens contactin 2 (axonal) (CNTN2), mRNA
NM_005116	Homo sapiens solute carrier family 23 (nucleobase transporters), member 1 (SLC23A1), mRNA
NM_005070	Homo sapiens solute carrier family 4, anion exchanger, member 3 (SLC4A3), mRNA
NM_005074	Homo sapiens solute carrier family 17 (sodium phosphate), member 1 (SLC17A1), mRNA
NM_005073	Homo sapiens solute carrier family 15 (oligopeptide transporter), member 1 (SLC15A1), mRNA
NM_005072	Homo sapiens solute carrier family 12 (potassium/chloride transporters), member 4 (SLC12A4), mRNA
NM 005063	Homo sapiens stearoyl-CoA desaturase (delta-9-desaturase) (SCD), mRNA
NM 005060	Homo sapiens RAR-related orphan receptor C (RORC), mRNA
NM 005059	Homo sapiens relaxin 2 (H2) (RLN2), mRNA
NM 005045	Homo sanjens reelin (RELN), mRNA
NM_005058	Homo sapiens RNA binding motif protein, Y chromosome, family 1, member A1 (RBMY1A1), mRNA
NM_005052	Homo sapiens ras-related C3 botulinum toxin substrate 3 (rho family, small GTP binding protein Rac3) (RAC3), mRNA
NM 005051	Homo sapiens glutaminyl-tRNA synthetase (QARS), mRNA
NM 005048	Homo sapiens parathyroid hormone receptor 2 (PTHR2), mRNA
NM 005044	Homo sapiens protein kinase, X-linked (PRKX), mRNA
NM 005043	Homo sapiens mitogen-activated protein kinase kinase 7 (MAP2K7), mRNA
NM 005042	Homo sapiens proline-rich protein HaeIII subfamily 2 (PRH2), mRNA
NM 005041	Homo sapiens perforin 1 (preforming protein) (PRF1), mRNA
NM 005040	Homo sapiens prolylcarboxypeptidase (angiotensinase C) (PRCP), mRNA
NM 005039	Homo sapiens proline-rich protein BstNI subfamily 1 (PRB1), mRNA
NM 005038	Homo sapiens peptidylprolyl isomerase D (cyclophilin D) (PPID), mRNA
NM_005029	Homo sapiens paired-like homeodomain transcription factor 3 (PITX3), mRNA
NM_005027	Homo sapiens phosphoinositide-3-kinase, regulatory subunit, polypeptide 2 (p85 beta) (PIK3R2), mRNA
NM_005026	Homo sapiens phosphoinositide-3-kinase, catalytic, delta polypeptide (PIK3CD), mRNA
NM_005021	Homo sapiens ectonucleotide pyrophosphatase/phosphodiesterase 3 (ENPP3), mRNA
NM 005019	Homo sapiens phosphodiesterase 1A, calmodulin-dependent (PDE1A), mRNA
NM 005018	Homo sapiens programmed cell death 1 (PDCD1), mRNA
NM 005015	Homo sapiens oxidase (cytochrome c) assembly 1-like (OXA1L), mRNA
NM_005085	Homo sapiens nucleoporin 214kD (CAIN) (NUP214), mRNA
NM 005124	Homo sapiens nucleoporin 153kD (NUP153), mRNA
NM 005013	Homo sapiens nucleobindin 2 (NUCB2), mRNA
NM 005012	Homo sapiens receptor tyrosine kinase-like orphan receptor 1 (ROR1), mRNA
NM 005011	Homo sapiens nuclear respiratory factor 1 (NRF1), mRNA
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27.5.005010	Homo sapiens neuronal cell adhesion molecule (NRCAM), mRNA
NM_005010	Homo sapiens non-metastatic cells 4, protein expressed in (NME4), mRNA
NM 005009	Homo sapiens nuclear factor of kappa light polypeptide gene enhancer in B-cells
NM_005007	inhibitor-like 1 (NFKBIL1), mRNA
> 7 5 005004	Homo sapiens NADH dehydrogenase (ubiquinone) 1 beta subcomplex, 8 (19kD,
NM_005004	Homo sapiens NADA denydrogenase (dordanione) i bear subcomplets, o (1712)
275 005001	ASHI) (NDUFB8), mRNA Homo sapiens NADH dehydrogenase (ubiquinone) 1 alpha subcomplex, 7
NM_005001	(14.5kD, B14.5a) (NDUFA7), mRNA
27.6.004000	Homo sapiens melanoma antigen, family A, 1 (directs expression of antigen
NM_004988	MZ2-E) (MAGEA1), mRNA
NR 005007	Homo sapiens leucine-rich, glioma inactivated 1 (LGI1), mRNA
NM_005097	Homo sapiens kinesin family member 5A (KIF5A), mRNA
NM_004984	Homo sapiens potassium inwardly-rectifying channel, subfamily J, member 9
NM_004983	(KCNJ9), mRNA
NTM 004092	Homo sapiens potassium inwardly-rectifying channel, subfamily J, member 8
NM_004982	(KCNJ8), mRNA
NM 000890	Homo sapiens potassium inwardly-rectifying channel, subfamily J, member 5
14141 000020	(KCNIS) mRNA
NM 004981	Homo sapiens potassium inwardly-rectifying channel, subfamily J, member 4
14141_004301	(KCNI4) mRNA
NM 005136	Homo sapiens potassium voltage-gated channel, Isk-related family, member 2
14141_005150	(KCNE2) mRNA
NM_004980	Homo sapiens potassium voltage-gated channel, Shal-related subfamily, member
1111_00-1500	3 (KCND3), mRNA
NM_004979	Homo sapiens potassium voltage-gated channel, Shal-related family, member 1
1117_00 1575	(KCNDI), mRNA
NM 004978	Homo sapiens potassium voltage-gated channel, Shaw-related subfamily,
2 12.2	member 4 (KCNC4), mRNA
NM_004977	Homo sapiens potassium voltage-gated channel, Shaw-related subfamily,
	member 3 (KCNC3), mRNA
NM_004976	Homo sapiens potassium voltage-gated channel, Shaw-related subfamily,
-	member 1 (KCNC1), mRNA
NM 004975	Homo sapiens potassium voltage-gated channel, Shab-related subfamily, member
	1 (KCNB1), mRNA
NM_004969	Homo sapiens insulin-degrading enzyme (IDE), mRNA
NM_005143	Homo sapiens haptoglobin (HP), mRNA
NM_004965	Homo sapiens high-mobility group (nonhistone chromosomal) protein 14
	(HMG14), mRNA
NM_005130	Homo sapiens heparin-binding growth factor binding protein (HBP17), mRNA
NM_004963	Homo sapiens guanylate cyclase 2C (heat stable enterotoxin receptor)
	(GUCY2C), mRNA
NM_005100	Homo sapiens A kinase (PRKA) anchor protein (gravin) 12 (AKAP12), mRNA
NM_005113	Homo sapiens golgi autoantigen, golgin subfamily a, 5 (GOLGA5), mRNA
NM_005145	Homo sapiens guanine nucleotide binding protein (G protein), gamma 7
77.5 0055 15	(GNG7), mRNA
NM_005142	Homo sapiens gastric intrinsic factor (vitamin B synthesis) (GIF), mRNA
NM 005110	Homo sapiens glutamine-fructose-6-phosphate transaminase 2 (GFPT2), mRNA
NM_004960	Homo sapiens fusion, derived from t(12;16) malignant liposarcoma (FUS), mRNA
NM 004959	Homo sapiens nuclear receptor subfamily 5, group A, member 1 (NR5A1),
1117_00,755	mRNA
NM 004957	Homo sapiens folylpolyglutamate synthase (FPGS), mRNA

NM_004956	Homo sapiens ets variant gene 1 (ETV1), mRNA
NM 004955	Homo sapiens solute carrier family 29 (nucleoside transporters), member 1
_	(SLC29A1), mRNA
NM 005107	Homo sapiens endonuclease G-like 1 (ENDOGL1), mRNA
NM 004953	Homo sapiens eukaryotic translation initiation factor 4 gamma, 1 (EIF4G1),
	mRNA
NM 004952	Homo sapiens ephrin-A3 (EFNA3), mRNA
NM 004944	Homo sapiens deoxyribonuclease I-like 3 (DNASE1L3), mRNA
NM 004938	Homo sapiens death-associated protein kinase 1 (DAPK1), mRNA
NM 005127	Homo saniens C-type (calcium dependent, carbohydrate-recognition domain)
	lectin, superfamily member 2 (activation-induced) (CLECSF2), mRNA
NM 004935	Homo sapiens cyclin-dependent kinase 5 (CDK5), mRNA
NM 004931	Homo sapiens CD8 antigen, beta polypeptide 1 (p37) (CD8B1), mRNA
NM 005125	Homo saniens copper chaperone for superoxide dismutase (CCS), mRNA
NM_005093	Homo sapiens core-binding factor, runt domain, alpha subunit 2; translocated to,
2.4.2_00000	2 (CBFA2T2), mRNA
NM 004930	Homo sapiens capping protein (actin filament) muscle Z-line, beta (CAPZB),
	mRNA
NM 005139	Homo sapiens annexin A3 (ANXA3), mRNA
NM 000664	Homo sapiens acetyl-Coenzyme A carboxylase alpha (ACACA), mRNA
NM 002108	Homo sapiens histidine ammonia-lyase (HAL), mRNA
NM 001718	Homo sapiens bone morphogenetic protein 6 (BMP6), mRNA
NM 001154	Homo sapiens annexin A5 (ANXA5), mRNA
NM 001153	Homo sapiens annexin A4 (ANXA4), mRNA
NM 004817	Homo sapiens tight junction protein 2 (zona occludens 2) (TJP2), mRNA
NM 004736	Homo sapiens xenotropic and polytropic retrovirus receptor (XPR1), mRNA
NM_004628	Homo sapiens xeroderma pigmentosum, complementation group C (XPC), mRNA
NM 004627	Homo sapiens tryptophan rich basic protein (WRB), mRNA
NM 004183	Homo sapiens vitelliform macular dystrophy (Best disease, bestrophin) (VMD2),
	mRNA
NM_004664	Homo sapiens Vertebrate LIN7 homolog 1, Tax interaction protein 33 (VELI1), mRNA
NM 004679	Homo sapiens variable charge, Y chromosome (VCY), mRNA
NM 004182	Homo sapiens ubiquitously-expressed transcript (UXT), mRNA
NM 004651	Homo sapiens ubiquitin specific protease 11 (USP11), mRNA
NM_004181	Homo sapiens ubiquitin carboxyl-terminal esterase L1 (ubiquitin thiolesterase) (UCHL1), mRNA
NM 004223	Homo sapiens ubiquitin-conjugating enzyme E2L 6 (UBE2L6), mRNA
NM 004223	Homo sapiens tetratricopeptide repeat domain 4 (TTC4), mRNA
NM 004622	Homo sapiens translin (TSN), mRNA
NM 004236	Homo sapiens thyroid receptor interacting protein 15 (TRIP15), mRNA
NM 004909	Homo sapiens taxol resistance associated gene 3 (TRAG3), mRNA
NM 004295	Homo sapiens TNF receptor-associated factor 4 (TRAF4), mRNA
NM 004179	Homo sapiens tryptophan hydroxylase (tryptophan 5-monooxygenase) (TPH),
	mRNA
NM_004195	Homo sapiens tumor necrosis factor receptor superfamily, member 18 (TNFRSF18), mRNA
NM 004202	Homo sapiens thymosin, beta 4, Y chromosome (TMSB4Y), mRNA
NM 004616	Homo sapiens transmembrane 4 superfamily member 3 (TM4SF3), mRNA
NM 004615	Homo sapiens transmembrane 4 superfamily member 2 (TM4SF2), mRNA
NM 004865	Homo sapiens TBP-like 1 (TBPL1), mRNA

NM_004613	Homo sapiens transglutaminase 2 (C polypeptide, protein-glutamine-gamma-glutamyltransferase) (TGM2), mRNA
	Homo sapiens transforming growth factor, beta receptor I (activin A receptor
NM_004612	type II-like kinase, 53kD) (TGFBR1), mRNA
NM 004708	Homo sapiens programmed cell death 5 (PDCD5), mRNA
NM 004918	Homo saniens T-cell leukemia/lymphoma 1B (TCL1B), mRNA
NM 004609	Homo saniens transcription factor 15 (basic helix-loop-helix) (TCF15), mRNA
NM 004780	Homo sapiens transcription elongation factor A (SII)-like 1 (TCEAL1), mRNA
	Homo sapiens thousand and one amino acid protein kinase (TAO1), mRNA
NM_004783	Homo sapiens TATA box binding protein (TBP)-associated factor, RNA
NM_004606	polymerase II, A, 250kD (TAF2A), mRNA
NM 004710	Homo sapiens synaptogyrin 2 (SYNGR2), mRNA
NM 004711	Homo saniens synaptogyrin 1 (SYNGR1), mRNA
NM_004605	Homo sapiens sulfotransferase family, cytosolic, 2B, member 1 (SULT2B1), mRNA
NM 004853	Homo sapiens syntaxin 8 (STX8), mRNA
NM 004603	Homo sapiens syntaxin 1A (brain) (STX1A), mRNA
	Homo sapiens serine/threonine kinase 12 (STK12), mRNA
NM_004217	Homo sapiens sterol regulatory element binding transcription factor 2 (SREBF2)
NM_004599	mRNA
NM_004176	Homo sapiens sterol regulatory element binding transcription factor 1 (SREBF1) mRNA
NM 000582	Homo sapiens secreted phosphoprotein 1 (osteopontin, bone sialoprotein I, early
111.12_00000	T-lymphocyte activation 1) (SPP1), mRNA
NM 004189	Homo sapiens SRY (sex determining region Y)-box 14 (SOX14), mRNA
	Homo sapiens small nuclear ribonucleoprotein polypeptide A (SNRPA), mRNA
NM_004596	Homo sapiens synaptosomal-associated protein, 29kD (SNAP29), mRNA
NM_004782	Homo sapiens synapiosomai-associated protein, 20kB (01/12/2), m2 4
NM_004595	Homo sapiens spermine synthase (SMS), mRNA
NM_004594	Homo sapiens solute carrier family 9 (sodium/hydrogen exchanger), isoform 5 (SLC9A5), mRNA
NM_004173	Homo sapiens solute carrier family 7 (cationic amino acid transporter, y+ system), member 4 (SLC7A4), mRNA
NM_004211	Homo sapiens solute carrier family 6 (neurotransmitter transporter, glycine),
NM_004858	Homo sapiens solute carrier family 4, sodium bicarbonate cotransporter, membe 8 (SLC4A8), mRNA
NM_004727	Homo sapiens solute carrier family 24 (sodium/potassium/calcium exchanger), member 1 (SLC24A1), mRNA
NM_004172	Homo sapiens solute carrier family 1 (glial high affinity glutamate transporter), member 3 (SLC1A3), nuclear gene encoding mitochondrial protein, mRNA
	Homo sapiens solute carrier family 1 (glial high affinity glutamate transporter),
NM_004171	member 2 (SLC1A2), nuclear gene encoding mitochondrial protein, mRNA
NM_004731	Homo sapiens solute carrier family 16 (monocarboxylic acid transporters),
NM_004695	Homo sapiens solute carrier family 16 (monocarboxylic acid transporters),
NM_004207	Homo sapiens solute carrier family 16 (monocarboxylic acid transporters), member 3 (SLC16A3), mRNA
NM 004870	Homo sapiens mannose-P-dolichol utilization defect 1 (MPDU1), mRNA
NM 004768	Homo sapiens splicing factor, arginine/serine-rich 11 (SFRS11), mRNA
	Homo sapiens sema domain, immunoglobulin domain (Ig), short basic domain,
NM_004636	secreted, (semaphorin) 3B (SEMA3B), mRNA

NM_004753	Homo sapiens short-chain dehydrogenase/reductase 1 (SDR1), mRNA
NM 004168	Homo sapiens succinate dehydrogenase complex, subunit A, flavoprotein (Fp)
14141_004100	(SDHA), nuclear gene encoding mitochondrial protein, mRNA
NM 004713	Homo sapiens serologically defined colon cancer antigen 1 (SDCCAG1), mRNA
NM_004591	Homo sapiens small inducible cytokine subfamily A (Cys-Cys), member 20
14141_004331	(SCYA20), mRNA
NM 004590	Homo sapiens small inducible cytokine subfamily A (Cys-Cys), member 16
14141_004230	(SCYA16), mRNA
ND 4 004500	Homo sapiens sodium channel, voltage-gated, type II, beta polypeptide
NM_004588	(SCN2B), mRNA
27.6.004165	Homo sapiens Ras-related associated with diabetes (RRAD), mRNA
NM_004165	Homo sapiens ribosomal protein S6 kinase, 90kD, polypeptide 5 (RPS6KA5),
NM_004755	l =
27.4.004506	mRNA Homo sapiens ribosomal protein S6 kinase, 90kD, polypeptide 3 (RPS6KA3),
NM_004586	
27.6.004700	mRNA Homo sapiens solute carrier family 22 (organic anion transporter), member 6
NM_004790	
27.5.004050	(SLC22A6), mRNA Homo sapiens RecQ protein-like 5 (RECQL5), mRNA
NM_004259	Homo sapiens RecQ protein-like 5 (RECQL5), iniciva
NM_004260	Homo sapiens RecQ protein-like 4 (RECQL4), mRNA
NM 004583	Homo sapiens RAB5C, member RAS oncogene family (RAB5C), mRNA
NM_004582	Homo sapiens Rab geranylgeranyltransferase, beta subunit (RABGGTB), mRNA
NM_004581	Homo sapiens Rab geranylgeranyltransferase, alpha subunit (RABGGTA),
	mRNA
NM_004251	Homo sapiens RAB9, member RAS oncogene family (RAB9), mRNA
NM_004162	Homo sapiens RAB5A, member RAS oncogene family (RAB5A), mRNA
NM_004578	Homo sapiens RAB4, member RAS oncogene family (RAB4), mRNA
NM_004914	Homo sapiens RAB36, member RAS oncogene family (RAB36), mRNA
NM_004580	Homo sapiens RAB27A, member RAS oncogene family (RAB27A), mRNA
NM_004663	Homo sapiens RAB11A, member RAS oncogene family (RAB11A), mRNA
NM_004160	Homo sapiens peptide YY (PYY), mRNA
NM_004103	Homo sapiens protein tyrosine kinase 2 beta (PTK2B), mRNA
NM_004158	Homo sapiens persephin (PSPN), mRNA
NM_004577	Homo sapiens phosphoserine phosphatase (PSPH), mRNA
NM_004159	Homo sapiens proteasome (prosome, macropain) subunit, beta type, 8 (large multifunctional protease 7) (PSMB8), mRNA
NM 004917	Homo sapiens kallikrein 4 (prostase, enamel matrix, prostate) (KLK4), mRNA
NM 004157	Homo sapiens protein kinase, cAMP-dependent, regulatory, type II, alpha
11112_00-115/	(PRKAR2A), mRNA
NM_004758	Homo sapiens peripheral benzodiazepine receptor-associated protein 1 (PRAX-
1111_004756	1), mRNA
NM_004576	Homo sapiens protein phosphatase 2 (formerly 2A), regulatory subunit B (PR
11171_004370	52), beta isoform (PPP2R2B), mRNA
NM 004156	Homo sapiens protein phosphatase 2 (formerly 2A), catalytic subunit, beta
11171_007130	isoform (PPP2CB), mRNA
NM_000942	Homo sapiens peptidylprolyl isomerase B (cyclophilin B) (PPIB), mRNA
NM 004575	Homo sapiens POU domain, class 4, transcription factor 2 (POU4F2), mRNA
NM 004573	Homo sapiens phospholipase C, beta 2 (PLCB2), mRNA
NM 004573	Homo sapiens plakophilin 2 (PKP2), mRNA
	Homo sapiens PBX/knotted 1 hoemobox 1 (PKNOX1), mRNA
NM 004571	Homo sapiens rembrane-associated tyrosine- and threonine-specific cdc2-
NM_004203	inhibitory kinase (PKMYT1), mRNA
NM_004910	Homo sapiens phosphatidylinositol transfer protein, membrane-associated
14141 004910	Tromo sapiens phosphandymiositor transfer protein, memorane-associated

	(PITPNM), mRNA
NM 004278	Homo sapiens phosphatidylinositol glycan, class L (PIGL), mRNA
NM_004569	Homo sapiens phosphatidylinositol glycan, class H (PIGH), mRNA
NM 004855	Homo sapiens phosphatidylinositol glycan, class B (PIGB), mRNA
NM_004862	Homo sapiens LPS-induced TNF-alpha factor (PIG7), mRNA
NM 004878	Homo sapiens prostaglandin E synthase (PTGES), mRNA
NM 004567	Homo sapiens 6-phosphofructo-2-kinase/fructose-2,6-biphosphatase 4
	(PFKFB4), mRNA
NM_004566	Homo sapiens 6-phosphofructo-2-kinase/fructose-2,6-biphosphatase 3 (PFKFB3), mRNA
NM_004836	Homo sapiens eukaryotic translation initiation factor 2-alpha kinase 3 (EIF2AK3), mRNA
NM_004716	Homo sapiens proprotein convertase subtilisin/kexin type 7 (PCSK7), mRNA
NM_000437	Homo sapiens platelet-activating factor acetylhydrolase 2 (40kD) (PAFAH2), mRNA
NM_004199	Homo sapiens procollagen-proline, 2-oxoglutarate 4-dioxygenase (proline 4-hydroxylase), alpha polypeptide II (P4HA2), mRNA
NM_004154	Homo sapiens pyrimidinergic receptor P2Y, G-protein coupled, 6 (P2RY6), mRNA
NM_004280	Homo sapiens eukaryotic translation elongation factor 1 epsilon 1 (EEF1E1), mRNA
NM_004741	Homo sapiens nucleolar phosphoprotein p130 (P130), mRNA
NM_004802	Homo sapiens otoferlin (OTOF), mRNA
NM_004852	Homo sapiens one cut domain, family member 2 (ONECUT2), mRNA
NM_004254	Homo sapiens solute carrier family 22 (organic anion transporter), member 8 (SLC22A8), mRNA
NM_004298	Homo sapiens nucleoporin 155kD (NUP155), mRNA
NM_004560	Homo sapiens receptor tyrosine kinase-like orphan receptor 2 (ROR2), mRNA
NM_004822	Homo sapiens netrin 1 (NTN1), mRNA
NM_004796	Homo sapiens neurexin 3 (NRXN3), mRNA
NM_004558	Homo sapiens neurturin (NRTN), mRNA
NM_004688	Homo sapiens N-myc (and STAT) interactor (NMI), mRNA
NM_004148	Homo sapiens ninjurin 1 (NINJ1), mRNA
NM_004552	Homo sapiens NADH dehydrogenase (ubiquinone) Fe-S protein 5 (15kD) (NADH-coenzyme Q reductase) (NDUFS5), mRNA
NM_004551	Homo sapiens NADH dehydrogenase (ubiquinone) Fe-S protein 3 (30kD) (NADH-coenzyme Q reductase) (NDUFS3), mRNA
NM_004550	Homo sapiens NADH dehydrogenase (ubiquinone) Fe-S protein 2 (49kD) (NADH-coenzyme Q reductase) (NDUFS2), mRNA
NM 004540	Homo sapiens neural cell adhesion molecule 2 (NCAM2), mRNA
NM_004644	Homo sapiens adaptor-related protein complex 3, beta 2 subunit (AP3B2), mRNA
NM_004538	Homo sapiens nucleosome assembly protein 1-like 3 (NAP1L3), mRNA
NM 004145	Homo sapiens myosin IXB (MYO9B), mRNA
NM_004294	Homo sapiens mitochondrial translational release factor 1 (MTRF1), mRNA
NM_004923	Homo sapiens metallothionein-like 5, testis-specific (tesmin) (MTL5), mRNA
NM_004143	Homo sapiens Cbp/p300-interacting transactivator, with Glu/Asp-rich carboxy-terminal domain, 1 (CITED1), mRNA
NM 004279	Homo sapiens pentidase (mitochondrial processing) beta (PMPCB), mRNA
	Homo sapiens molybdenum cofactor synthesis 2 (MOCS2), mRNA
NM 004528	Homo sapiens microsomal glutathione S-transferase 3 (MGST3), mRNA
NM 004279 NM 004531 NM 004244	Homo sapiens peptidase (mitochondrial processing) beta (PMPCB), mRNA Homo sapiens molybdenum cofactor synthesis 2 (MOCS2), mRNA Homo sapiens CD163 antigen (CD163), mRNA
14141_004528	rionio sapiens microsomai giutatnione o-transferase o (MOS10), micha

Homo sapiens MFH-amplified sequences with leucine-rich tandem repeats 1 (MASL1), mRNA
Homo sapiens mannosidase, alpha, class 2A, member 1 (MAN2A1), mRNA
Homo sapiens mitogen-activated protein kinase kinase kinase 13 (MAP3K13), mRNA
Homo sapiens low density lipoprotein-related protein 1 (alpha-2-macroglobulin receptor) (LRP1), mRNA
Homo sapiens protease, serine, 15 (PRSS15), mRNA
Homo sapiens LIM homeobox protein 2 (LHX2), mRNA
Homo sapiens serine palmitoyltransferase, long chain base subunit 2 (SPTLC2), mRNA
Homo sapiens like-glycosyltransferase (LARGE), mRNA
Homo sapiens klotho (KL), mRNA
Homo sapiens kinesin family member 5B (KIF5B), mRNA
Homo sapiens kinesin heavy chain member 2 (KIF2), mRNA
Homo sapiens apoptosis-associated tyrosine kinase (AATK), mRNA
Homo sapiens potassium voltage-gated channel, KQT-like subfamily, member 4 (KCNO4), mRNA
Homo sapiens potassium voltage-gated channel, KQT-like subfamily, member 3 (KCNQ3), mRNA
Homo sapiens potassium voltage-gated channel, KQT-like subfamily, member 2 (KCNQ2), mRNA
Homo sapiens potassium large conductance calcium-activated channel, subfamily M, beta member 1 (KCNMB1), mRNA
Homo sapiens potassium voltage-gated channel, shaker-related subfamily, beta member 3 (KCNAB3), mRNA
Homo sapiens cytokeratin type II (K6HF), mRNA
Homo sapiens integrin, beta-like 1 (with EGF-like repeat domains) (ITGBL1), mRNA
Homo sapiens integrin-linked kinase (ILK), mRNA
Homo sapiens interleukin enhancer binding factor 1 (ILF1), mRNA
Homo sapiens interleukin 1 receptor, type II (IL1R2), mRNA
Homo sapiens interleukin 16 (lymphocyte chemoattractant factor) (IL16), mRNA
Homo sapiens interleukin 11 receptor, alpha (IL11RA), mRNA
Homo sapiens immunoglobulin superfamily, member 2 (IGSF2), mRNA
Homo sapiens isocitrate dehydrogenase 3 (NAD+) gamma (IDH3G), mRNA
Homo sapiens heat shock 70kD protein 9B (mortalin-2) (HSPA9B), mRNA
Homo sapiens PRP4/STK/WD splicing factor (HPRP4P), mRNA
Homo sapiens U4/U6-associated RNA splicing factor (HPRP3P), mRNA
Homo sapiens homeo box C6 (HOXC6), mRNA
Homo sapiens homeo box B7 (HOXB7), mRNA
Homo sapiens hepatocyte nuclear factor 3, gamma (HNF3G), mRNA
Homo sapiens hepatocyte nuclear factor 3, alpha (HNF3A), mRNA
Homo sapiens hepatocyte growth factor-regulated tyrosine kinase substrate (HGS), mRNA
Homo sapiens mitogen-activated protein kinase kinase kinase kinase 4 (MAP4K4), mRNA
Homo sapiens hepatoma-derived growth factor (high-mobility group protein 1-
like) (HDGF), mRNA Homo sapiens zinc finger protein 254 (ZNF254), mRNA

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NM 004893 Ho NM 004130 Ho NM 004286 Ho NM 004128 Ho (CO NM 004491 Ho NM 004890 Ho NM 004810 Ho NM 004224 Ho NM 004871 Ho NM 004487 Ho Trail	tomo sapiens H2A histone family, member Y (H2AFY), mRNA tomo sapiens glycogenin (GYG), mRNA tomo sapiens GTP binding protein 1 (GTPBP1), mRNA tomo sapiens general transcription factor IIF, polypeptide 2 (30kD subunit) GTF2F2), mRNA tomo sapiens glucocorticoid receptor DNA binding factor 1 (GRLF1), mRNA tomo sapiens glutamate receptor, ionotropic, AMPA 2 (GRIA2), mRNA tomo sapiens growth factor receptor-bound protein 14 (GRB14), mRNA tomo sapiens GRB2-related adaptor protein 2 (GRAP2), mRNA tomo sapiens G protein-coupled receptor 50 (GPR50), mRNA tomo sapiens golgi SNAP receptor complex member 1 (GOSR1), mRNA tomo sapiens golgi autoantigen, golgin subfamily b, macrogolgin (with ransmembrane signal), 1 (GOLGB1), mRNA tomo sapiens guanine nucleotide binding protein 11 (GNG11), mRNA tomo sapiens guanine nucleotide binding protein (G protein), alpha 14 GNA14), mRNA tomo sapiens glucagon-like peptide 2 receptor (GLP2R), mRNA
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NM_004224 Hd NM_004871 Hd NM_004487 Hd tra	Iomo sapiens G protein-coupled receptor 50 (GPR50), mRNA Iomo sapiens golgi SNAP receptor complex member 1 (GOSR1), mRNA Iomo sapiens golgi autoantigen, golgin subfamily b, macrogolgin (with ransmembrane signal), 1 (GOLGB1), mRNA Iomo sapiens guanine nucleotide binding protein 11 (GNG11), mRNA Iomo sapiens guanine nucleotide binding protein (G protein), alpha 14 GNA14), mRNA Iomo sapiens glucagon-like peptide 2 receptor (GLP2R), mRNA
NM_004871 He NM_004487 He tra	Iomo sapiens golgi SNAP receptor complex member 1 (GOSR1), mRNA Iomo sapiens golgi autoantigen, golgin subfamily b, macrogolgin (with ransmembrane signal), 1 (GOLGB1), mRNA Iomo sapiens guanine nucleotide binding protein 11 (GNG11), mRNA Iomo sapiens guanine nucleotide binding protein (G protein), alpha 14 GNA14), mRNA Iomo sapiens glucagon-like peptide 2 receptor (GLP2R), mRNA
NM_004487 Ho	ansmembrane signal), 1 (GOLGB1), mRNA Iomo sapiens guanine nucleotide binding protein 11 (GNG11), mRNA Iomo sapiens guanine nucleotide binding protein (G protein), alpha 14 GNA14), mRNA Iomo sapiens glucagon-like peptide 2 receptor (GLP2R), mRNA
	Iomo sapiens guanine nucleotide binding protein 11 (GNG11), mRNA Iomo sapiens guanine nucleotide binding protein (G protein), alpha 14 GNA14), mRNA Iomo sapiens glucagon-like peptide 2 receptor (GLP2R), mRNA
NM 004126 H	Iomo sapiens guanine nucleotide binding protein (G protein), alpha 14 GNA14), mRNA Iomo sapiens glucagon-like peptide 2 receptor (GLP2R), mRNA
	GNA14), mRNA Iomo sapiens glucagon-like peptide 2 receptor (GLP2R), mRNA
NM_004297 H	Iomo sapiens glucagon-like peptide 2 receptor (GLP2R), mRNA
(0	Iomo sapiens glucagon-like peptide 2 receptor (GLP2R), mRNA
NM_004246 H	Iomo saniens gastric inhibitary nolymentide (GIP) mPNA
	Iomo sapiens gamma-glutamyltransferase-like activity 1 (GGTLA1), mRNA
	Iomo sapiens geranylgeranyl diphosphate synthase 1 (GGPS1), mRNA
NM_004188 H	Iomo sapiens growth factor independent 1B (potential regulator of CDKN1A,
	anslocated in CML) (GFI1B), mRNA
	Iomo sapiens guanine deaminase (GDA), mRNA
	Iomo sapiens glucosaminyl (N-acetyl) transferase 3, mucin type (GCNT3), nRNA
NM 004193 H	Iomo sapiens golgi-specific brefeldin A resistance factor 1 (GBF1), mRNA
NM 002030 H	Iomo sapiens formyl peptide receptor-like 2 (FPRL2), mRNA
NM_004476 H	Iomo sapiens folate hydrolase (prostate-specific membrane antigen) 1 (FOLH1), nRNA
	Iomo sapiens fms-related tyrosine kinase 3 (FLT3), mRNA
	Iomo sapiens flotillin 2 (FLOT2), mRNA
	Iomo sapiens forkhead box D1 (FOXD1), mRNA
	Iomo sapiens forkhead box G1A (FOXG1A), mRNA
	Iomo sapiens forkhead box D2 (FOXD2), mRNA
	Iomo sapiens c-fos induced growth factor (vascular endothelial growth factor D)
	FIGF), mRNA
	Iomo sapiens four and a half LIM domains 3 (FHL3), mRNA
	Iomo sapiens farnesyl-diphosphate farnesyltransferase 1 (FDFT1), mRNA
	Iomo sapiens Fc fragment of IgG, receptor, transporter, alpha (FCGRT), mRNA
	Iomo sapiens fatty acid synthase (FASN), mRNA
	Iomo sapiens phenylalanine-tRNA synthetase-like (FARSL), mRNA
	Homo sapiens coagulation factor II (thrombin) receptor-like 2 (F2RL2), mRNA
	Homo sapiens Kruppel-like factor 4 (gut) (KLF4), mRNA
NM_004455 H	Homo sapiens exostoses (multiple)-like 1 (EXTL1), mRNA
	Iomo sapiens ets variant gene 5 (ets-related molecule) (ETV5), mRNA
	Iomo sapiens electron-transferring-flavoprotein dehydrogenase (ETFDH),
	nuclear gene encoding mitochondrial protein, mRNA
	Iomo sapiens estrogen-related receptor beta (ESRRB), mRNA
	Homo sapiens protein disulfide isomerase related protein (calcium-binding
	protein, intestinal-related) (ERP70), mRNA
NM_004911 H	Iomo sapiens protein disulfide isomerase related protein (calcium-binding

	Homo sapiens epidermal growth factor receptor pathway substrate 8 (EPS8),
NM_004447	mRNA
NM 004446	Homo sapiens glutamyl-prolyl-tRNA synthetase (EPRS), mRNA
NM 004431	Homo sapiens EphA2 (EPHA2), mRNA
NM_004099	Homo sapiens erythrocyte membrane protein band 7.2 (stomatin) (EPB72),
NM_004437	Homo sapiens erythrocyte membrane protein band 4.1 (elliptocytosis 1, RH-linked) (EPR41), mRNA
NM_004435	Homo sapiens endonuclease G (ENDOG), nuclear gene encoding mitochondrial
NM_004434	Homo sapiens echinoderm microtubule-associated protein-like (EMAPL),
NM_004433	Homo sapiens E74-like factor 3 (ets domain transcription factor, epithelial-
NM_004096	Homo sapiens eukaryotic translation initiation factor 4E binding protein 2 (FIF4ERP2), mRNA
NM_004095	Homo sapiens eukaryotic translation initiation factor 4E binding protein 1 (EIF4EBP1), mRNA
NM 004430	Homo sapiens early growth response 3 (EGR3), mRNA
NM 004093	Homo sapiens ephrin-B2 (EFNB2), mRNA
NM 004429	Homo sapiens ephrin-B1 (EFNB1), mRNA
NM 004428	Homo sapiens ephrin-A1 (EFNA1), mRNA
	Homo sapiens integral membrane protein 2A (ITM2A), mRNA
NM_004867	Homo sapiens desmoplakin (DPI, DPII) (DSP), mRNA
NM_004415	Homo sapiens desinoplakin (Dr.), Dr. 11 (Dos.), indivis
NM_004760	Homo sapiens serine/threonine kinase 17a (apoptosis-inducing) (STK17A), mRNA
NM_004413	Homo sapiens dipeptidase 1 (renal) (DPEP1), mRNA
NM 004088	Homo sapiens deoxynucleotidyltransferase, terminal (DNTT), mRNA
NM 004412	Homo sapiens DNA (cytosine-5-)-methyltransferase 2 (DNMT2), mRNA
NM 004411	Homo sapiens dynein, cytoplasmic, intermediate polypeptide 1 (DNCI1), mRNA
NM 004407	Homo sapiens dentin matrix acidic phosphoprotein (DMP1), mRNA
NM_004746	Homo sapiens discs, large (Drosophila) homolog-associated protein 1 (DLGAP1), mRNA
NM_004747	Homo sapiens discs, large (Drosophila) homolog 5 (DLG5), mRNA
NM 004087	Homo sapiens discs, large (Drosophila) homolog 1 (DLG1), mRNA
NM_004900	Homo sapiens phorbolin (similar to apolipoprotein B mRNA editing protein) (DJ742C19.2), mRNA
NM_004404	Homo sapiens neural precursor cell expressed, developmentally down-regulated 5 (NEDD5), mRNA
NM_004402	Homo sapiens DNA fragmentation factor, 40 kD, beta polypeptide (caspase-activated DNase) (DFFB), mRNA
NM_004401	Homo sapiens DNA fragmentation factor, 45 kD, alpha polypeptide (DFFA), mRNA
NM_004083	Homo sapiens DNA-damage-inducible transcript 3 (DDIT3), mRNA
NM 004734	Homo sapiens doublecortin and CaM kinase-like 1 (DCAMKL1), mRNA
NM 004394	Homo sapiens death-associated protein (DAP), mRNA
NM_004393	Homo sapiens dystroglycan 1 (dystrophin-associated glycoprotein 1) (DAG1), mRNA
NM_004229	Homo sapiens cofactor required for Sp1 transcriptional activation, subunit 2 (150kD) (CRSP2), mRNA
NM 004079	Homo sapiens cathepsin S (CTSS), mRNA
NM 004390	Homo sapiens cathepsin H (CTSH), mRNA
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NM_004388	Homo sapiens chitobiase, di-N-acetyl- (CTBS), mRNA
NM_004387	Homo sapiens cardiac-specific homeo box (CSX), mRNA
NM_004861	Homo sapiens cerebroside (3'-phosphoadenylylsulfate:galactosylceramide 3')
	sulfotransferase (CST), mRNA
NM_004078	Homo sapiens cysteine and glycine-rich protein 1 (CSRP1), mRNA
NM_004386	Homo sapiens chondroitin sulfate proteoglycan 3 (neurocan) (CSPG3), mRNA
NM_004385	Homo sapiens chondroitin sulfate proteoglycan 2 (versican) (CSPG2), mRNA
NM_004384	Homo sapiens casein kinase 1, gamma 3 (CSNK1G3), mRNA
NM_004383	Homo sapiens c-src tyrosine kinase (CSK), mRNA
NM_004075	Homo sapiens cryptochrome 1 (photolyase-like) (CRY1), mRNA
NM 004778	Homo sapiens G protein-coupled receptor 44 (GPR44), mRNA
NM_004750	Homo sapiens cytokine receptor-like factor 1 (CRLF1), mRNA
NM_004382	Homo sapiens corticotropin releasing hormone receptor 1 (CRHR1), mRNA
NM_004379	Homo sapiens cAMP responsive element binding protein 1 (CREB1), mRNA
NM_004377	Homo sapiens carnitine palmitoyltransferase I, muscle (CPT1B), mRNA
NM_004748	Homo sapiens cell cycle progression 8 protein (CPR8), mRNA
NM_004074	Homo sapiens cytochrome c oxidase subunit VIII (COX8), nuclear gene
	encoding mitochondrial protein, mRNA
NM_004766	Homo sapiens coatomer protein complex, subunit beta 2 (beta prime) (COPB2), mRNA
NM_004645	Homo sapiens coilin (COIL), mRNA
NM 000614	Homo sapiens ciliary neurotrophic factor (CNTF), mRNA
NM 004368	Homo sapiens calponin 2 (CNN2), mRNA
NM 004072	Homo sapiens chemokine-like receptor 1 (CMKLR1), mRNA
NM 004071	Homo sapiens CDC-like kinase1 (CLK1), mRNA
NM 004362	Homo sapiens calmegin (CLGN), mRNA
NM_004070	Homo sapiens chloride channel Ka (CLCNKA), mRNA
NM_004804	Homo sapiens WD40 protein Ciao1 (CIAO1), mRNA
NM_004267	Homo sapiens carbohydrate (chondroitin 6/keratan) sulfotransferase 2 (CHST2), mRNA
NM 004067	Homo sapiens chimerin (chimaerin) 2 (CHN2), mRNA
NM_004284	Homo sapiens chromodomain helicase DNA binding protein 1-like (CHD1L), mRNA
NM_004364	Homo sapiens CCAAT/enhancer binding protein (C/EBP), alpha (CEBPA), mRNA
NM 004065	Homo sapiens cerebellar degeneration-related protein (34kD) (CDR1), mRNA
NM_004233	Homo sapiens CD83 antigen (activated B lymphocytes, immunoglobulin
NR 004056	superfamily) (CD83), mRNA
NM_004356	Homo sapiens CD81 antigen (target of antiproliferative antibody 1) (CD81), mRNA
NM_004357	Homo sapiens CD151 antigen (CD151), mRNA
NM_004350	Homo sapiens runt-related transcription factor 3 (RUNX3), mRNA
NM_004349	Homo sapiens core-binding factor, runt domain, alpha subunit 2; translocated to, 1; cyclin D-related (CBFA2T1), mRNA
NM 004345	Homo sapiens cathelicidin antimicrobial peptide (CAMP), mRNA
NM_000722	Homo sapiens calcium channel, voltage-dependent, alpha 2/delta subunit 1 (CACNA2D1), mRNA
NM_004334	Homo sapiens bone marrow stromal cell antigen 1 (BST1), mRNA
NM_004887	Homo sapiens small inducible cytokine subfamily B (Cys-X-Cys), member 14 (BRAK) (SCYB14), mRNA
NM_004333	Homo sapiens v-raf murine sarcoma viral oncogene homolog B1 (BRAF), mRNA

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NM_004329	Homo sapiens bone morphogenetic protein receptor, type IA (BMPR1A), mRNA
NM_004827	Homo sapiens ATP-binding cassette, sub-family G (WHITE), member 2
	(ABCG2), mRNA
NM_004326	Homo sapiens B-cell CLL/lymphoma 9 (BCL9), mRNA
NM_004765	Homo sapiens B-cell CLL/lymphoma 7C (BCL7C), mRNA
NM 004324	Homo sapiens BCL2-associated X protein (BAX), mRNA
NM_004656	Homo sapiens BRCA1 associated protein-1 (ubiquitin carboxy-terminal
_	hydrolase) (BAP1), mRNA
NM 004048	Homo sapiens beta-2-microglobulin (B2M), mRNA
NM 004655	Homo sapiens axin 2 (conductin, axil) (AXIN2), mRNA
NM 004321	Homo sapiens axonal transport of synaptic vesicles (ATSV), mRNA
NM 004888	Homo sapiens ATPase, H+ transporting, lysosomal (vacuolar proton pump),
_	member J (ATP6J), mRNA
NM_004047	Homo sapiens ATPase, H+ transporting, lysosomal (vacuolar proton pump)
<u> </u>	21kD (ATP6F), mRNA
NM_004046	Homo sapiens ATP synthase, H+ transporting, mitochondrial F1 complex, alpha
_	subunit, isoform 1, cardiac muscle (ATP5A1), mRNA
NM 001683	Homo sapiens ATPase, Ca++ transporting, plasma membrane 2 (ATP2B2),
_	mRNA
NM 004314	Homo sapiens ADP-ribosyltransferase 1 (ART1), mRNA
NM 004313	Homo sapiens arrestin, beta 2 (ARRB2), mRNA
NM 004312	Homo sapiens arrestin 3, retinal (X-arrestin) (ARR3), mRNA
NM 004311	Homo sapiens ADP-ribosylation factor-like 3 (ARL3), mRNA
NM 004675	Homo sapiens ras homolog gene family, member I (ARHI), mRNA
NM 004310	Homo sapiens ras homolog gene family, member H (ARHH), mRNA
NM 004309	Homo sapiens Rho GDP dissociation inhibitor (GDI) alpha (ARHGDIA), mRNA
NM 004308	Homo sapiens Rho GTPase activating protein 1 (ARHGAP1), mRNA
NM 004040	Homo sapiens ras homolog gene family, member B (ARHB), mRNA
NM 004290	Homo sapiens ring finger protein 14 (RNF14), mRNA
NM 004797	Homo sapiens adipose most abundant gene transcript 1 (APM1), mRNA
NM 004039	Homo sapiens annexin A2 (ANXA2), mRNA
NM 004306	Homo sapiens annexin A13 (ANXA13), mRNA
NM 004038	Homo sapiens amylase, alpha 1A; salivary (AMY1A), mRNA
NM 004305	Homo sapiens bridging integrator 1 (BIN1), mRNA
NM 004857	Homo sapiens A kinase (PRKA) anchor protein 5 (AKAP5), mRNA
NM 004833	Homo sapiens absent in melanoma 2 (AIM2), mRNA
NM 004208	Homo sapiens programmed cell death 8 (apoptosis-inducing factor) (PDCD8),
1111_004200	mRNA
NM 002199	Homo sapiens interferon regulatory factor 2 (IRF2), mRNA
NM 001569	Homo sapiens interleukin-1 receptor-associated kinase 1 (IRAK1), mRNA
NM 001567	Homo sapiens inositol polyphosphate phosphatase-like 1 (INPPL1), mRNA
NM_002194	Homo sapiens inositol polyphosphate-1-phosphatase (INPP1), mRNA
NM 002111	Homo sapiens huntingtin (Huntington disease) (HD), mRNA
NM 000165	Homo sapiens muntingtin (Huntington disease) (HD), mkNA Homo sapiens gap junction protein, alpha 1, 43kD (connexin 43) (GJA1), mRNA
NM 001999	Homo sapiens fibrillin 2 (congenital contractural arachnodactyly) (FBN2),
1001999	mRNA
NM 001937	Homo sapiens dermatopontin (DPT), mRNA
NM 001381	Homo sapiens docking protein 1, 62kD (downstream of tyrosine kinase 1)
	(DOK1), mRNA
NM 000729	Homo sapiens cholecystokinin (CCK), mRNA
NM_000486	Homo sapiens aquaporin 2 (coll cting duct) (AQP2), mRNA
NM 001520	Homo sapiens general transcription factor IIIC, polypeptide 1 (alpha subunit,
TATAT OOTOTO	I monto sapiens general transcription factor mic, poryperhide I (aipha subunit,

	220kD) (GTF3C1), mRNA
NM_002097	Homo sapiens general transcription factor IIIA (GTF3A), mRNA
NM 003205	Homo sapiens transcription factor 12 (HTF4, helix-loop-helix transcription
	factors 4) (TCF12), mRNA
NM_000440	Homo sapiens phosphodiesterase 6A, cGMP-specific, rod, alpha (PDE6A),
	mRNA
NM_000806	Homo sapiens gamma-aminobutyric acid (GABA) A receptor, alpha 1
_	(GABRA1), mRNA
NM_001809	Homo sapiens centromere protein A (17kD) (CENPA), mRNA
NM 000439	Homo sapiens proprotein convertase subtilisin/kexin type 1 (PCSK1), mRNA
NM_002529	Homo sapiens neurotrophic tyrosine kinase, receptor, type 1 (NTRK1), mRNA
NM_003417	Homo sapiens zinc finger protein 264 (ZNF264), mRNA
NM_000395	Homo sapiens colony stimulating factor 2 receptor, beta, low-affinity
	(granulocyte-macrophage) (CSF2RB), mRNA
NM_000065	Homo sapiens complement component 6 (C6), mRNA
NM_000252	Homo sapiens myotubular myopathy 1 (MTM1), mRNA
NM_000229	Homo sapiens lecithin-cholesterol acyltransferase (LCAT), nuclear gene
	encoding mitochondrial protein, mRNA
NM_000224	Homo sapiens keratin 18 (KRT18), mRNA
NM_000211	Homo sapiens integrin, beta 2 (antigen CD18 (p95), lymphocyte function-
	associated antigen 1; macrophage antigen 1 (mac-1) beta subunit) (ITGB2),
	mRNA
NM_000208	Homo sapiens insulin receptor (INSR), mRNA
NM_000206	Homo sapiens interleukin 2 receptor, gamma (severe combined
	immunodeficiency) (IL2RG), mRNA
NM_000416	Homo sapiens interferon gamma receptor 1 (IFNGR1), mRNA
NM_000201	Homo sapiens intercellular adhesion molecule 1 (CD54), human rhinovirus
	receptor (ICAM1), mRNA
NM_000350	Homo sapiens ATP-binding cassette, sub-family A (ABC1), member 4
	(ABCA4), mRNA
NM_000110	Homo sapiens dihydropyrimidine dehydrogenase (DPYD), mRNA
NM_000375	Homo sapiens uroporphyrinogen III synthase (congenital erythropoietic
	porphyria) (UROS), mRNA
NM_000459	Homo sapiens TEK tyrosine kinase, endothelial (venous malformations, multiple
27.5.004.050	cutaneous and mucosal) (TEK), mRNA
NM_001053	Homo sapiens somatostatin receptor 5 (SSTR5), mRNA
NM_001052	Homo sapiens somatostatin receptor 4 (SSTR4), mRNA
NM_001051	Homo sapiens somatostatin receptor 3 (SSTR3), mRNA
NM_001050	Homo sapiens somatostatin receptor 2 (SSTR2), mRNA
NM_001049	Homo sapiens somatostatin receptor 1 (SSTR1), mRNA
NM_000348	Homo sapiens steroid-5-alpha-reductase, alpha polypeptide 2 (3-oxo-5 alpha-
ND (000040	steroid delta 4-dehydrogenase alpha 2) (SRD5A2), mRNA
NM_000340	Homo sapiens solute carrier family 2 (facilitated glucose transporter), member 2
NR4 000220	(SLC2A2), mRNA
NM_000338	Homo sapiens solute carrier family 12 (sodium/potassium/chloride transporters),
NR4 000001	member 1 (SLC12A1), mRNA
NM_000231	Homo sapiens sarcoglycan, gamma (35kD dystrophin-associated glycoprotein)
NTM 001024	(SGCG), mRNA
NM_001034	Homo sapiens ribonucleotide reductase M2 polypeptide (RRM2), mRNA
NM_000448	Homo sapiens recombination activating gene 1 (RAG1), mRNA
NM_000303	Homo sapiens phosphomannomutase 2 (PMM2), mRNA
NM_000302	Homo sapiens procollagen-lysine, 2-oxoglutarate 5-dioxygenase (lysine

	hydroxylase, Ehlers-Danlos syndrome type VI) (PLOD), mRNA
NM_000282	Homo sapiens propionyl Coenzyme A carboxylase, alpha polypeptide (PCCA),
NIVI_000262	nuclear gene encoding mitochondrial protein, mRNA
ND 6 000001	Homo sapiens 6-pyruvoyl-tetrahydropterin synthase/dimerization cofactor of
NM_000281	hepatocyte nuclear factor 1 alpha (TCF1) (PCBD), mRNA
>D (000077	nepatocyte nuclear factor i atpina (1011) (1019), nuclear
NM_000277	Homo sapiens phenylalanine hydroxylase (PAH), mRNA
NM_000436	Homo sapiens 3-oxoacid CoA transferase (OXCT), nuclear gene encoding mitochondrial protein, mRNA
NM_000274	Homo sapiens ornithine aminotransferase (gyrate atrophy) (OAT), nuclear gene encoding mitochondrial protein, mRNA
NM 000273	Homo sapiens ocular albinism 1 (Nettleship-Falls) (OA1), mRNA
NM 000272	Homo sapiens nephronophthisis 1 (juvenile) (NPHP1), mRNA
NM 000271	Homo sapiens Niemann-Pick disease, type C1 (NPC1), mRNA
NM_000269	Homo sapiens non-metastatic cells 1, protein (NM23A) expressed in (NME1), mRNA
NM 000268	Homo sapiens neurofibromin 2 (bilateral acoustic neuroma) (NF2), mRNA
NM_000267	Homo sapiens neurofibromin 1 (neurofibromatosis, von Recklinghausen disease, Watson disease) (NF1), mRNA
NM 000434	Homo sapiens sialidase 1 (lysosomal sialidase) (NEU1), mRNA
	Homo sapiens Norrie disease (pseudoglioma) (NDP), mRNA
NM_000266	Homo sapiens Norrie disease (pseudognoma) (NDI), intervi-
NM_000265	disease, autosomal 1) (NCF1), mRNA
NM_000262	Homo sapiens N-acetylgalactosaminidase, alpha- (NAGA), mRNA
NM_000261	Homo sapiens myocilin, trabecular meshwork inducible glucocorticoid response (MYOC), mRNA
NM_000258	Homo sapiens myosin, light polypeptide 3, alkali; ventricular, skeletal, slow (MYL3), mRNA
NM_000432	Homo sapiens myosin, light polypeptide 2, regulatory, cardiac, slow (MYL2), mRNA
NM_000257	Homo sapiens myosin, heavy polypeptide 7, cardiac muscle, beta (MYH7), mRNA
NM_000431	Homo sapiens mevalonate kinase (mevalonic aciduria) (MVK), mRNA
NM_000255	Homo sapiens methylmalonyl Coenzyme A mutase (MUT), nuclear gene
	encoding mitochondrial protein, mRNA
NM_000254	Homo sapiens 5-methyltetrahydrofolate-homocysteine methyltransferase (MTR), mRNA
NM_000253	Homo sapiens microsomal triglyceride transfer protein (large polypeptide, 88kD) (MTP), mRNA
NM_000250	Homo sapiens myeloperoxidase (MPO), nuclear gene encoding mitochondrial protein, mRNA
NM 000248	Homo sapiens microphthalmia-associated transcription factor (MITF), mRNA
NM 000247	Homo sapiens MHC class I polypeptide-related sequence A (MICA), mRNA
NM 000246	Homo sapiens MHC class II transactivator (MHC2TA), mRNA
NM_000245	Homo sapiens met proto-oncogene (hepatocyte growth factor receptor) (MET), mRNA
NM_000244	Homo sapiens multiple endocrine neoplasia I (MEN1), mRNA
NM 000243	Homo sapiens Mediterranean fever (MEFV), mRNA
	Homo sapiens mannose-binding lectin (protein C) 2, soluble (opsonic defect)
NM_000242	(MBL2), mRNA
NM_000429	Homo sapiens methionine adenosyltransferase I, alpha (MAT1A), mRNA
NM_000240	Homo sapiens monoamine oxidase A (MAOA), nuclear gene encoding mitochondrial protein, mRNA

NM_000428	Homo sapiens latent transforming growth factor beta binding protein 2 (LTBP2), mRNA
NM_000238	Homo sapiens potassium voltage-gated channel, subfamily H (eag-related), member 2 (KCNH2), mRNA
NM 000237	Homo sapiens lipoprotein lipase (LPL), mRNA
NM 000427	Homo sapiens loricrin (LOR), mRNA
NM 000236	Homo saniens lipase, hepatic (LIPC), mRNA
NM_000235	Homo sapiens lipase A, lysosomal acid, cholesterol esterase (Wolman disease) (LIPA), mRNA
NM 000234	Homo sapiens ligase I, DNA, ATP-dependent (LIG1), mRNA
NM_000233	Homo sapiens luteinizing hormone/choriogonadotropin receptor (LHCGR), mRNA
NM_000228	Homo sapiens laminin, beta 3 (nicein (125kD), kalinin (140kD), BM600 (125kD)) (LAMB3), mRNA
NM_000426	Homo sapiens laminin, alpha 2 (merosin, congenital muscular dystrophy) (I.AMA2), mRNA
NM_000226	Homo sapiens keratin 9 (epidermolytic palmoplantar keratoderma) (KRT9), mRNA
NM_000422	Homo sapiens keratin 17 (KRT17), mRNA
NM_000223	Homo sapiens keratin 12 (Meesmann corneal dystrophy) (KRT12), mRNA
NM_000421	Homo sapiens keratin 10 (epidermolytic hyperkeratosis; keratosis palmaris et plantaris) (KRT10), mRNA
NM_000222	Homo sapiens v-kit Hardy-Zuckerman 4 feline sarcoma viral oncogene homolog (KIT), mRNA
NM_000218	Homo sapiens potassium voltage-gated channel, KQT-like subfamily, member 1 (KCNQ1), mRNA
NM_000219	Homo sapiens potassium voltage-gated channel, Isk-related family, member 1 (KCNE1), mRNA
NM_000217	Homo sapiens potassium voltage-gated channel, shaker-related subfamily, member 1 (episodic ataxia with myokymia) (KCNA1), mRNA
NM_000216	Homo sapiens Kallmann syndrome 1 sequence (KAL1), mRNA
NM_000215	Homo sapiens Janus kinase 3 (a protein tyrosine kinase, leukocyte) (JAK3), mRNA
NM_000212	Homo sapiens integrin, beta 3 (platelet glycoprotein IIIa, antigen CD61) (ITGB3), mRNA
NM_000209	Homo sapiens insulin promoter factor 1, homeodomain transcription factor (IPF1), mRNA
NM 000207	Homo sapiens insulin (INS), mRNA
NM 000418	Homo sapiens interleukin 4 receptor (ILAR), mRNA
NM_000417	Homo sapiens interleukin 2 receptor, alpha (IL2RA), mRNA
NM_001551	Homo sapiens immunoglobulin (CD79A) binding protein 1 (IGBP1), mRNA
NM_000203	Homo sapiens iduronidase, alpha-L- (IDUA), mRNA
NM_000415	Homo sapiens islet amyloid polypeptide (IAPP), mRNA
NM_000200	Homo sapiens histatin 3 (HTN3), mRNA
NM_001538	Homo sapiens heat shock transcription factor 4 (HSF4), mRNA
NM_000859	Homo sapiens 3-hydroxy-3-methylglutaryl-Coenzyme A reductase (HMGCR), mRNA
NM_001527	Homo sapiens histone deacetylase 2 (HDAC2), mRNA
NM_001525	Homo sapiens hypocretin (orexin) receptor 1 (HCRTR1), mRNA
NM_001524	Homo sapiens hypocretin (orexin) neuropeptide precursor (HCRT), mRNA
NM_001510	Homo sapiens glutamate receptor, ionotropic, delta 2 (GRID2), mRNA
NM_000829	Homo sapiens glutamate receptor, ionotrophic, AMPA 4 (GRIA4), mRNA

NM_001496	Homo sapiens GDNF family receptor alpha 3 (GFRA3), mRNA
NM 001486	Homo sapiens glucokinase (hexokinase 4) regulatory protein (GCKR), mRNA
NM 000820	Homo sapiens growth arrest-specific 6 (GAS6), mRNA
NM 000155	Homo sapiens galactose-1-phosphate uridylyltransferase (GALT), mRNA
NM 000153	Homo sapiens galactosylceramidase (Krabbe disease) (GALC), mRNA
NM 000816	Homo sapiens gamma-aminobutyric acid (GABA) A receptor, gamma 2
	(GABRG2), mRNA
NM 000815	Homo sapiens gamma-aminobutyric acid (GABA) A receptor, delta (GABRD),
112.2_00000	mRNA
NM_000811	Homo sapiens gamma-aminobutyric acid (GABA) A receptor, alpha 6
1442_000011	(GABRA6), mRNA
NM 000809	Homo sapiens gamma-aminobutyric acid (GABA) A receptor, alpha 4
112.12_00000	(GABRA4), mRNA
NM 000808	Homo sapiens gamma-aminobutyric acid (GABA) A receptor, alpha 3
1442_00000	(GABRA3), mRNA
NM_000807	Homo sapiens gamma-aminobutyric acid (GABA) A receptor, alpha 2
11112_00000	(GABRA2), mRNA
NM 000151	Homo sapiens glucose-6-phosphatase, catalytic (glycogen storage disease type I,
	von Gierke disease) (G6PC), mRNA
NM 001452	Homo sapiens forkhead box F2 (FOXF2), mRNA
NM 000138	Homo sapiens fibrillin 1 (Marfan syndrome) (FBN1), mRNA
NM 000136	Homo sapiens Fanconi anemia, complementation group C (FANCC), mRNA
NM 001445	Homo sapiens fatty acid binding protein 6, ileal (gastrotropin) (FABP6), mRNA
NM 001442	Homo sapiens fatty acid binding protein 4, adipocyte (FABP4), mRNA
NM 001443	Homo sapiens fatty acid binding protein 1, liver (FABP1), mRNA
NM 001441	Homo sapiens fatty acid amide hydrolase (FAAH), mRNA
NM 000401	Homo sapiens exostoses (multiple) 2 (EXT2), mRNA
NM 000127	Homo sapiens exostoses (multiple) 1 (EXT1), mRNA
NM 001433	Homo sapiens ER to nucleus signalling 1 (ERN1), mRNA
NM 000122	Homo sapiens excision repair cross-complementing rodent repair deficiency,
1,111_000122	complementation group 3 (xeroderma pigmentosum group B complementing)
	(ERCC3), mRNA
NM 000121	Homo sapiens erythropoietin receptor (EPOR), mRNA
NM 000120	Homo sapiens epoxide hydrolase 1, microsomal (xenobiotic) (EPHX1), mRNA
NM 000119	Homo sapiens erythrocyte membrane protein band 4.2 (EPB42), mRNA
NM 001429	Homo sapiens E1A binding protein p300 (EP300), mRNA
NM 000118	Homo sapiens endoglin (Osler-Rendu-Weber syndrome 1) (ENG), mRNA
NM_000117	Homo sapiens emerin (Emery-Dreifuss muscular dystrophy) (EMD), mRNA
NM_001422	Homo sapiens E74-like factor 5 (ets domain transcription factor) (ELF5), mRNA
NM 000114	Homo sapiens endothelin 3 (EDN3), mRNA
NM_001393	Homo sapiens extracellular matrix protein 2, female organ and adipocyte specific
1111_001050	(ECM2), mRNA
NM 000112	Homo sapiens solute carrier family 26 (sulfate transporter), member 2
11112_000112	(SLC26A2), mRNA
NM_001382	Homo sapiens dolichyl-phosphate (UDP-N-acetylglucosamine) N-
1411_001502	acetylglucosaminephosphotransferase 1 (GlcNAc-1-P transferase) (DPAGT1),
	mRNA
NM 001365	Homo sapiens discs, large (Drosophila) homolog 4 (DLG4), mRNA
NM 000792	Homo sapiens deiodinase, iodothyronine, type I (DIO1), mRNA
NM 001358	Homo sapiens DEAD/H (Asp-Glu-Ala-Asp/His) box polypeptide 15 (DDX15),
14141_001338	mRNA
NM 000107	Homo sapiens damage-specific DNA binding protein 2 (48kD) (DDB2), mRNA
141AT 00010\	1 Homo sapiens damage-specific DIVA uniting protein 2 (40kD) (DDB2), likeWA

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NM_001348	Homo sapiens death-associated protein kinase 3 (DAPK3), mRNA
NM_000101	Homo sapiens cytochrome b-245, alpha polypeptide (CYBA), mRNA
NM 001081	Homo sapiens cubilin (intrinsic factor-cobalamin receptor) (CUBN), mRNA
NM 001334	Homo sapiens cathepsin O (CTSO), mRNA
NM 001328	Homo sapiens C-terminal binding protein 1 (CTBP1), mRNA
NM 000554	Homo sapiens cone-rod homeobox (CRX), mRNA
NM 000096	Homo sapiens ceruloplasmin (ferroxidase) (CP), mRNA
NM_000095	Homo sapiens cartilage oligomeric matrix protein (pseudoachondroplasia,
	eniphyseal dysplasia 1, multiple) (COMP), mRNA
NM_000392	Homo sapiens ATP-binding cassette, sub-family C (CFTR/MRP), member 2
1111_0000	(ABCC2), mRNA
NM 000085	Homo sapiens chloride channel Kb (CLCNKB), mRNA
NM 000084	Homo sapiens chloride channel 5 (nephrolithiasis 2, X-linked, Dent disease)
	(CLCN5), mRNA
NM 001279	Homo sapiens cell death-inducing DFFA-like effector a (CIDEA), mRNA
NM_000080	Homo sapiens cholinergic receptor, nicotinic, epsilon polypeptide (CHRNE),
	mRNA
NM 000751	Homo sapiens cholinergic receptor, nicotinic, delta polypeptide (CHRND),
	mRNA
NM 000747	Homo sapiens cholinergic receptor, nicotinic, beta polypeptide 1 (muscle)
	(CHRNB1), mRNA
NM 000079	Homo sapiens cholinergic receptor, nicotinic, alpha polypeptide 1 (muscle)
	(CHRNA1), mRNA
NM 001273	Homo sapiens chromodomain helicase DNA binding protein 4 (CHD4), mRNA
NM 001271	Homo sapiens chromodomain helicase DNA binding protein 2 (CHD2), mRNA
NM 001270	Homo sapiens chromodomain helicase DNA binding protein 1 (CHD1), mRNA
NM 000078	Homo sapiens cholesteryl ester transfer protein, plasma (CETP), mRNA
NM 000076	Homo sapiens cyclin-dependent kinase inhibitor 1C (p57, Kip2) (CDKN1C),
11112_00001	mRNA
NM 001258	Homo sapiens cyclin-dependent kinase 3 (CDK3), mRNA
NM 001251	Homo sapiens CD68 antigen (CD68), mRNA
NM 000074	Homo sapiens tumor necrosis factor (ligand) superfamily, member 5 (hyper-IgM
	syndrome) (TNFSF5), mRNA
NM_000073	Homo sapiens CD3G antigen, gamma polypeptide (TiT3 complex) (CD3G),
1111_000075	mRNA
NM 001249	Homo sapiens ectonucleoside triphosphate diphosphohydrolase 5 (ENTPD5),
11112_002215	mRNA
NM_001248	Homo sapiens ectonucleoside triphosphate diphosphohydrolase 3 (ENTPD3),
11112_001210	mRNA
NM 001246	Homo sapiens ectonucleoside triphosphate diphosphohydrolase 2 (ENTPD2),
11212_001210	mRNA
NM 000072	Homo sapiens CD36 antigen (collagen type I receptor, thrombospondin receptor)
	(CD36), mRNA
NM_000591	Homo sapiens CD14 antigen (CD14), mRNA
NM 000071	Homo sapiens cystathionine-beta-synthase (CBS), mRNA
NM 000388	Homo sapiens calcium-sensing receptor (hypocalciuric hypercalcemia 1, severe
11117_000000	neonatal hyperparathyroidism) (CASR), mRNA
NM 000070	Homo sapiens calpain 3, (p94) (CAPN3), mRNA
NM_000069	Homo sapiens calcium channel, voltage-dependent, L type, alpha 1S subunit
14147_000009	(CACNA1S), mRNA
NM_001215	Homo sapiens carbonic anhydrase VI (CA6), mRNA
NM 000067	Homo sapiens carbonic anhydrase II (CA2), mRNA
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NM 000606	Homo sapiens complement component 8, gamma polypeptide (C8G), mRNA
NM 000066	Homo sapiens complement component 8, beta polypeptide (C8B), mRNA
NM 000562	Homo sapiens complement component 8, alpha polypeptide (C8A), mRNA
NM 000587	Homo sapiens complement component 7 (C7), mRNA
NM 000064	Homo sapiens complement component 3 (C3), mRNA
NM 000061	Homo sapiens Bruton agammaglobulinemia tyrosine kinase (BTK), mRNA
NM 001206	Homo sapiens basic transcription element binding protein 1 (BTEB1), mRNA
NM 000060	Homo sapiens biotinidase (BTD), mRNA
NM 001201	Homo sapiens bone morphogenetic protein 3 (osteogenic) (BMP3), mRNA
NM 001200	Homo sapiens bone morphogenetic protein 2 (BMP2), mRNA
NM 000386	Homo sapiens bleomycin hydrolase (BLMH), mRNA
NM 000057	Homo sapiens Bloom syndrome (BLM), mRNA
NM 001198	Homo sapiens PR domain containing 1, with ZNF domain (PRDM1), mRNA
NM 001196	Homo sapiens BH3 interacting domain death agonist (BID), mRNA
NM 000056	Homo sapiens branched chain keto acid dehydrogenase E1, beta polypeptide
	(maple syrup urine disease) (BCKDHB), nuclear gene encoding mitochondrial
	protein, mRNA
NM 000465	Homo sapiens BRCA1 associated RING domain 1 (BARD1), mRNA
NM 000705	Homo sapiens ATPase, H+/K+ exchanging, beta polypeptide (ATP4B), mRNA
NM 000049	Homo sapiens aspartoacylase (aminoacylase 2, Canavan disease) (ASPA),
_	mRNA
NM 000046	Homo sapiens arylsulfatase B (ARSB), mRNA
NM 000639	Homo sapiens tumor necrosis factor (ligand) superfamily, member 6 (TNFSF6),
_	mRNA
NM_000042	Homo sapiens apolipoprotein H (beta-2-glycoprotein I) (APOH), mRNA
NM 000041	Homo sapiens apolipoprotein E (APOE), mRNA
NM 000040	Homo sapiens apolipoprotein C-III (APOC3), mRNA
NM 000039	Homo sapiens apolipoprotein A-I (APOA1), mRNA
NM_000038	Homo sapiens adenomatosis polyposis coli (APC), mRNA
NM_001157	Homo sapiens annexin A11 (ANXA11), mRNA
NM_001147	Homo sapiens angiopoietin 2 (ANGPT2), mRNA
NM_001145	Homo sapiens angiogenin, ribonuclease, RNase A family, 5 (ANG), mRNA
NM_000036	Homo sapiens adenosine monophosphate deaminase 1 (isoform M) (AMPD1),
	mRNA
NM_001141	Homo sapiens arachidonate 15-lipoxygenase, second type (ALOX15B), mRNA
NM_000035	Homo sapiens aldolase B, fructose-bisphosphate (ALDOB), mRNA
NM_000034	Homo sapiens aldolase A, fructose-bisphosphate (ALDOA), mRNA
NM_000032	Homo sapiens aminolevulinate, delta-, synthase 2 (sideroblastic/hypochromic
	anemia) (ALAS2), nuclear gene encoding mitochondrial protein, mRNA
NM_000030	Homo sapiens alanine-glyoxylate aminotransferase (oxalosis I; hyperoxaluria I;
	glycolicaciduria; serine-pyruvate aminotransferase) (AGXT), mRNA
NM_001126	Homo sapiens adenylosuccinate synthase (ADSS), mRNA
NM_000684	Homo sapiens adrenergic, beta-1-, receptor (ADRB1), mRNA
NM_001125	Homo sapiens ADP-ribosylarginine hydrolase (ADPRH), mRNA
NM_001116	Homo sapiens adenylate cyclase 9 (ADCY9), mRNA
NM_001115	Homo sapiens adenylate cyclase 8 (brain) (ADCY8), mRNA
NM_001114	Homo sapiens adenylate cyclase 7 (ADCY7), mRNA
NM_001109	Homo sapiens a disintegrin and metalloproteinase domain 8 (ADAM8), mRNA
NM_001110	Homo sapiens a disintegrin and metalloproteinase domain 10 (ADAM10),
	mRNA (1.07m2) PNA
NM_001108	Homo sapiens acylphosphatase 2, muscle type (ACYP2), mRNA
NM_001107	Homo sapiens acylphosphatase 1, erythrocyte (common) type (ACYP1), mRNA

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NM_001104	Homo sapiens actinin, alpha 3 (ACTN3), mRNA
NM_001086	Homo sapiens arylacetamide deacetylase (esterase) (AADAC), mRNA
NM_001043	Homo sapiens solute carrier family 6 (neurotransmitter transporter,
	noradrenalin), member 2 (SLC6A2), mRNA
NM_000532	Homo sapiens propionyl Coenzyme A carboxylase, beta polypeptide (PCCB),
	nuclear gene encoding mitochondrial protein, mRNA
NM_002579	Homo sapiens paralemmin (PALM), mRNA
NM_002443	Homo sapiens microseminoprotein, beta- (MSMB), mRNA
NM_002418	Homo sapiens motilin (MLN), mRNA
NM_002300	Homo sapiens lactate dehydrogenase B (LDHB), mRNA
NM_002243	Homo sapiens potassium inwardly-rectifying channel, subfamily J, member 15
	(KCNJ15), mRNA
NM_001534	Homo sapiens homeo box 11-like 1 (HOX11L1), mRNA
NM_001454	Homo sapiens forkhead box J1 (FOXJ1), mRNA
NM_004001	Homo sapiens Fc fragment of IgG, low affinity IIb, receptor for (CD32) (FCGR2B), mRNA
NM 001276	Homo sapiens chitinase 3-like 1 (cartilage glycoprotein-39) (CHI3L1), mRNA
NM_001752	Homo sapiens catalase (CAT), mRNA
NM 001610	Homo sapiens acid phosphatase 2, lysosomal (ACP2), mRNA
NM 003461	Homo sapiens zyxin (ZYX), mRNA
NM 003460	Homo sapiens zona pellucida glycoprotein 2 (sperm receptor) (ZP2), mRNA
NM_003459	Homo sapiens solute carrier family 30 (zinc transporter), member 3 (SLC30A3),
1411_003 133	mRNA
NM 003430	Homo sapiens zinc finger protein 91 (HPF7, HTF10) (ZNF91), mRNA
NM 003429	Homo sapiens zinc finger protein 85 (HPF4, HTF1) (ZNF85), mRNA
NM 003428	Homo sapiens zinc finger protein 84 (HPF2) (ZNF84), mRNA
NM 003416	Homo sapiens zinc finger protein 7 (KOX 4, clone HF.16) (ZNF7), mRNA
NM 003427	Homo sapiens zinc finger protein 76 (expressed in testis) (ZNF76), mRNA
NM 003426	Homo sapiens zinc finger protein 74 (Cos52) (ZNF74), mRNA
NM_003425	Homo sapiens zinc finger protein 45 (a Kruppel-associated box (KRAB) domain
	polypeptide) (ZNF45), mRNA
NM 003423	Homo sapiens zinc finger protein 43 (HTF6) (ZNF43), mRNA
NM 003422	Homo sapiens zinc finger protein 42 (myeloid-specific retinoic acid-responsive)
	(ZNF42), mRNA
NM_003420	Homo sapiens zinc finger protein 35 (clone HF.10) (ZNF35), mRNA
NM_003458	Homo sapiens bassoon (presynaptic cytomatrix protein) (BSN), mRNA
NM_003456	Homo sapiens zinc finger protein 205 (ZNF205), mRNA
NM_003453	Homo sapiens zinc finger protein 198 (ZNF198), mRNA
NM 003450	Homo sapiens zinc finger protein 174 (ZNF174), mRNA
NM_003447	Homo sapiens zinc finger protein 165 (ZNF165), mRNA
NM_003446	Homo sapiens zinc finger protein 157 (HZF22) (ZNF157), mRNA
NM_003443	Homo sapiens zinc finger protein 151 (pHZ-67) (ZNF151), mRNA
NM_003442	Homo sapiens zinc finger protein 143 (clone pHZ-1) (ZNF143), mRNA
NM_003441	Homo sapiens zinc finger protein 141 (clone pHZ-44) (ZNF141), mRNA
NM_003440	Homo sapiens zinc finger protein 140 (clone pHZ-39) (ZNF140), mRNA
NM 003438	Homo sapiens zinc finger protein 137 (clone pHZ-30) (ZNF137), mRNA
NM_003437	Homo sapiens zinc finger protein 136 (clone pHZ-20) (ZNF136), mRNA
NM_003436	Homo sapiens zinc finger protein 135 (clone pHZ-17) (ZNF135), mRNA
NM_003435	Homo sapiens zinc finger protein 134 (clone pHZ-15) (ZNF134), mRNA
NM_003434	Homo sapiens zinc finger protein 133 (clone pHZ-13) (ZNF133), mRNA
NM_003433	Homo sapi ns zinc finger protein 132 (clone pHZ-12) (ZNF132), mRNA
NM_003431	Homo sapiens zinc finger protein 124 (HZF-16) (ZNF124), mRNA

NM_003411	Homo sapiens zinc finger protein, Y-linked (ZFY), mRNA
NM_003410	Homo sapiens zinc finger protein, X-linked (ZFX), mRNA
NM_003405	Homo sapiens tyrosine 3-monooxygenase/tryptophan 5-monooxygenase
	activation protein, eta polypeptide (YWHAH), mRNA
NM_003404	Homo sapiens tyrosine 3-monooxygenase/tryptophan 5-monooxygenase
	activation protein, beta polypeptide (YWHAB), mRNA
NM_000380	Homo sapiens xeroderma pigmentosum, complementation group A (XPA),
	mRNA
NM_003931	Homo sapiens WAS protein family, member 1 (WASF1), mRNA
NM_003384	Homo sapiens vaccinia related kinase 1 (VRK1), mRNA
NM_003383	Homo sapiens very low density lipoprotein receptor (VLDLR), mRNA
NM_003382	Homo sapiens vasoactive intestinal peptide receptor 2 (VIPR2), mRNA
NM_003381	Homo sapiens vasoactive intestinal peptide (VIP), mRNA
NM_003380	Homo sapiens vimentin (VIM), mRNA
NM_003377	Homo sapiens vascular endothelial growth factor B (VEGFB), mRNA
NM_003376	Homo sapiens vascular endothelial growth factor (VEGF), mRNA
NM_000376	Homo sapiens vitamin D (1,25- dihydroxyvitamin D3) receptor (VDR), mRNA
NM_003375	Homo sapiens voltage-dependent anion channel 2 (VDAC2), mRNA
NM_003374	Homo sapiens voltage-dependent anion channel 1 (VDAC1), mRNA
NM_003371	Homo sapiens vav 2 oncogene (VAV2), mRNA
NM_003370	Homo sapiens vasodilator-stimulated phosphoprotein (VASP), mRNA
NM_003762	Homo sapiens vesicle-associated membrane protein 4 (VAMP4), mRNA
NM_003369	Homo sapiens UV radiation resistance associated gene (UVRAG), mRNA
NM_003577	Homo sapiens undifferentiated embryonic cell transcription factor 1 (UTF1),
	mRNA
NM_003470	Homo sapiens ubiquitin specific protease 7 (herpes virus-associated) (USP7),
	mRNA
NM_003481	Homo sapiens ubiquitin specific protease 5 (isopeptidase T) (USP5), mRNA
NM_003363	Homo sapiens ubiquitin specific protease 4 (proto-oncogene) (USP4), mRNA
NM_003368	Homo sapiens ubiquitin specific protease 1 (USP1), mRNA
NM_003940	Homo sapiens ubiquitin specific protease 13 (isopeptidase T-3) (USP13), mRNA
NM_003367	Homo sapiens upstream transcription factor 2, c-fos interacting (USF2), mRNA
NM_003366	Homo sapiens ubiquinol-cytochrome c reductase core protein II (UQCRC2),
77.5.00005	mRNA
NM_003365	Homo sapiens ubiquinol-cytochrome c reductase core protein I (UQCRC1),
377 C 0022 C4	mRNA
NM_003364	Homo sapiens uridine phosphorylase (UP), mRNA
NM_003361	Homo sapiens uromodulin (uromucoid, Tamm-Horsfall glycoprotein) (UMOD), mRNA
NM 003709	Homo sapiens Kruppel-like factor 7 (ubiquitous) (KLF7), mRNA
NM 003709	Homo sapiens UDP glycosyltransferase 8 (UDP-galactose ceramide
14147_002200	galactosyltransferase) (UGT8), mRNA
NM 001074	Homo sapiens UDP glycosyltransferase 2 family, polypeptide B7 (UGT2B7),
14141_0010/4	mRNA
NM 001077	Homo sapiens UDP glycosyltransferase 2 family, polypeptide B17 (UGT2B17),
1414_0010//	mRNA
NM 001076	Homo sapiens UDP glycosyltransferase 2 family, polypeptide B15 (UGT2B15),
14141_001070	mRNA
NM 001075	Homo sapiens UDP glycosyltransferase 2 family, polypeptide B10 (UGT2B10),
1111_0010/3	mRNA
NM 003359	Homo sapiens UDP-glucose dehydrogenase (UGDH), mRNA
NM 003358	Homo sapiens UDP-glucose ceramide glucosyltransferase (UGCG), mRNA
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NM_003357	Homo sapiens uteroglobin (UGB), mRNA
NM_003352	Homo sapiens ubiquitin-like 1 (sentrin) (UBL1), mRNA
NM_003347	Homo sapiens ubiquitin-conjugating enzyme E2L 3 (UBE2L3), mRNA
NM_003337	Homo sapiens ubiquitin-conjugating enzyme E2B (RAD6 homolog) (UBE2B), mRNA
NM 003336	Homo sapiens ubiquitin-conjugating enzyme E2A (RAD6 homolog) (UBE2A),
MM_003330	mRNA
NM 003335	Homo sapiens ubiquitin-activating enzyme E1-like (UBE1L), mRNA
NM 000550	Homo sapiens tyrosinase-related protein 1 (TYRP1), mRNA
NM 000372	Homo sapiens tyrosinase (oculocutaneous albinism IA) (TYR), mRNA
NM 001071	Homo sapiens thymidylate synthetase (TYMS), mRNA
NM 003331	Homo sapiens tyrosine kinase 2 (TYK2), mRNA
NM 003330	Homo sapiens thioredoxin reductase 1 (TXNRD1), mRNA
NM 003329	Homo sapiens thioredoxin (TXN), mRNA
NM 003328	Homo sapiens TXK tyrosine kinase (TXK), mRNA
NM 003324	Homo sapiens tubby like protein 3 (TULP3), mRNA
NM 003323	Homo sapiens tubby like protein 2 (TULP2), mRNA
NM 003321	Homo sapiens Tu translation elongation factor, mitochondrial (TUFM), mRNA
NM 001070	Homo sapiens tubulin, gamma 1 (TUBG1), mRNA
NM 001069	Homo sapiens tubulin, beta polypeptide (TUBB), mRNA
NM 000371	Homo sapiens transthyretin (prealbumin, amyloidosis type I) (TTR), mRNA
NM 000370	Homo sapiens tocopherol (alpha) transfer protein (ataxia (Friedreich-like) with
14141_000370	vitamin E deficiency) (TTPA), mRNA
NM 003319	Homo sapiens titin (TTN), mRNA
NM 003318	Homo sapiens TTK protein kinase (TTK), mRNA
NM 003317	Homo sapiens thyroid transcription factor 1 (TITF1), mRNA
NM 003315	Homo sapiens tetratricopeptide repeat domain 2 (TTC2), mRNA
NM 003314	Homo sapiens tetratricopeptide repeat domain 1 (TTC1), mRNA
NM 003311	Homo sapiens tumor suppressing subtransferable candidate 3 (TSSC3), mRNA
NM 003310	Homo sapiens tumor suppressing subtransferable candidate 1 (TSSC1), mRNA
NM 000369	Homo sapiens thyroid stimulating hormone receptor (TSHR), mRNA
NM 000549	Homo sapiens thyroid stimulating hormone, beta (TSHB), mRNA
NM 003496	Homo sapiens transformation/transcription domain-associated protein (TRRAP),
NW_003490	mRNA
NM_003301	Homo sapiens thyrotropin-releasing hormone receptor (TRHR), mRNA
NM_003299	Homo sapiens tumor rejection antigen (gp96) 1 (TRA1), mRNA
NM_003298	Homo sapiens nuclear receptor subfamily 2, group C, member 2 (NR2C2), mRNA
NM 003296	Homo sapiens testis specific protein 1 (probe H4-1 p3-1) (TPX1), mRNA
NM 003295	Homo sapiens tumor protein, translationally-controlled 1 (TPT1), mRNA
NM 003595	Homo sapiens tyrosylprotein sulfotransferase 2 (TPST2), mRNA
NM_003292	Homo sapiens translocated promoter region (to activated MET oncogene) (TPR),
	mRNA
NM 003291	Homo sapiens tripeptidyl peptidase II (TPP2), mRNA
NM_000547	Homo sapiens thyroid peroxidase (TPO), nuclear gene encoding mitochondrial
	protein, mRNA
NM 003290	Homo sapiens tropomyosin 4 (TPM4), mRNA
NM 003289	Homo sapiens tropomyosin 2 (beta) (TPM2), mRNA
NM 000366	Homo sapiens tropomyosin 1 (alpha) (TPM1), mRNA
NM_000365	Homo sapiens triosephosphate isomerase 1 (TPI1), mRNA
NM_003288	Homo sapiens tumor protein D52-like 2 (TPD52L2), mRNA
NM 003288	Homo sapiens tumor protein D52-like 1 (TPD52L1), mRNA
14141 003201	110mo sapiens tunior protein D32-nao 1 (11 D32D1), metara

NM 003935	Homo sapiens topoisomerase (DNA) III beta (TOP3B), mRNA
NM 001067	Homo sapiens topoisomerase (DNA) II alpha (170kD) (TOP2A), mRNA
NM 003285	Homo sapiens tenascin R (restrictin, janusin) (TNR), mRNA
NM 003284	Homo sapiens transition protein 1 (during histone to protamine replacement)
	(TNP1), mRNA
NM 000364	Homo sapiens troponin T2, cardiac (TNNT2), mRNA
NM 003283	Homo sapiens troponin T1, skeletal, slow (TNNT1), mRNA
NM 000363	Homo sapiens troponin I, cardiac (TNNI3), mRNA
NM 003282	Homo sapiens troponin I, skeletal, fast (TNNI2), mRNA
NM 003281	Homo sapiens troponin I, skeletal, slow (TNNI1), mRNA
NM 003279	Homo sapiens troponin C2, fast (TNNC2), mRNA
NM 003280	Homo sapiens troponin C, slow (TNNC1), mRNA
NM 003985	Homo sapiens tyrosine kinase, non-receptor, 1 (TNK1), mRNA
NM_001244	Homo sapiens tumor necrosis factor (ligand) superfamily, member 8 (TNFSF8),
	mRNA
NM_001252	Homo sapiens tumor necrosis factor (ligand) superfamily, member 7 (TNFSF7), mRNA
NM_003326	Homo sapiens tumor necrosis factor (ligand) superfamily, member 4 (tax-transcriptionally activated glycoprotein 1, 34kD) (TNFSF4), mRNA
NTN / 002909	Homo sapiens tumor necrosis factor (ligand) superfamily, member 13
NM_003808	(TNFSF13), mRNA
NM_003809	Homo sapiens tumor necrosis factor (ligand) superfamily, member 12 (TNFSF12), mRNA
NM_003810	Homo sapiens tumor necrosis factor (ligand) superfamily, member 10 (TNFSF10), mRNA
NM_001243	Homo sapiens tumor necrosis factor receptor superfamily, member 8 (TNFRSF8), mRNA
NM_001242	Homo sapiens tumor necrosis factor receptor superfamily, member 7 (TNFRSF7), mRNA
NM_000043	Homo sapiens tumor necrosis factor receptor superfamily, member 6 (TNFRSF6), mRNA
NM_003327	Homo sapiens tumor necrosis factor receptor superfamily, member 4 (TNFRSF4), mRNA
NM_001066	Homo sapiens tumor necrosis factor receptor superfamily, member 1B (TNFRSF1B), mRNA
NM_001065	Homo sapiens tumor necrosis factor receptor superfamily, member 1A (TNFRSF1A), mRNA
NM_001192	Homo sapiens tumor necrosis factor receptor superfamily, member 17 (TNFRSF17), mRNA
NM_003820	Homo sapiens tumor necrosis factor receptor superfamily, member 14 (herpesvirus entry mediator) (TNFRSF14), mRNA
NM_003790	Homo sapiens tumor necrosis factor receptor superfamily, member 12 (translocating chain-association membrane protein) (TNFRSF12), mRNA
NM_002546	Homo sapiens tumor necrosis factor receptor superfamily, member 11b (osteoprotegerin) (TNFRSF11B), mRNA
NM_003839	Homo sapiens tumor necrosis factor receptor superfamily, member 11a, activator of NFKB (TNFRSF11A), mRNA
NM_003840	Homo sapiens tumor necrosis factor receptor superfamily, member 10d, decoy with truncated death domain (TNFRSF10D), mRNA
NM_003842	Homo sapiens tumor necrosis factor receptor superfamily, member 10b (TNFRSF10B), mRNA
NM 003844	Homo sapiens tumor necrosis factor receptor superfamily, member 10a

	(TNFRSF10A), mRNA
NM 003276	Homo sapiens thymopoietin (TMPO), mRNA
NM 003275	Homo sapiens tropomodulin (TMOD), mRNA
NM 003274	Homo sapiens transmembrane protein 1 (TMEM1), mRNA
NM 003692	Homo sapiens transmembrane protein with EGF-like and two follistatin-like
11112_00000	domains 1 (TMEFF1), mRNA
NM 003273	Homo sapiens transmembrane 7 superfamily member 2 (TM7SF2), mRNA
NM_003272	Homo sapiens transmembrane 7 superfamily member 1 (upregulated in kidney)
_	(TM7SF1), mRNA
NM 003271	Homo sapiens transmembrane 4 superfamily member 7 (TM4SF7), mRNA
NM 003270	Homo sapiens transmembrane 4 superfamily member 6 (TM4SF6), mRNA
NM 003963	Homo sapiens transmembrane 4 superfamily member 5 (TM4SF5), mRNA
NM_003269	Homo sapiens nuclear receptor subfamily 2, group E, member 1 (NR2E1),
	mRNA
NM_003266	Homo sapiens toll-like receptor 4 (TLR4), mRNA
NM_003265	Homo sapiens toll-like receptor 3 (TLR3), mRNA
NM_003264	Homo sapiens toll-like receptor 2 (TLR2), mRNA
NM_003263	Homo sapiens toll-like receptor 1 (TLR1), mRNA
NM_003258	Homo sapiens thymidine kinase 1, soluble (TK1), mRNA
NM_003257	Homo sapiens tight junction protein 1 (zona occludens 1) (TJP1), mRNA
NM_003256	Homo sapiens tissue inhibitor of metalloproteinase 4 (TIMP4), mRNA
NM_003254	Homo sapiens tissue inhibitor of metalloproteinase 1 (erythroid potentiating
	activity, collagenase inhibitor) (TIMP1), mRNA
NM_003597	Homo sapiens TGFB inducible early growth response 2 (TIEG2), mRNA
NM_003253	Homo sapiens T-cell lymphoma invasion and metastasis 1 (TIAM1), mRNA
NM_000460	Homo sapiens thrombopoietin (myeloproliferative leukemia virus oncogene
	ligand, megakaryocyte growth and development factor) (THPO), mRNA
NM_003249	Homo sapiens thimet oligopeptidase 1 (THOP1), mRNA
NM_003248	Homo sapiens thrombospondin 4 (THBS4), mRNA
NM_003247	Homo sapiens thrombospondin 2 (THBS2), mRNA
NM_003246	Homo sapiens thrombospondin 1 (THBS1), mRNA
NM_000361	Homo sapiens thrombomodulin (THBD), mRNA
NM_000360	Homo sapiens tyrosine hydroxylase (TH), mRNA
NM_003241	Homo sapiens transglutaminase 4 (prostate) (TGM4), mRNA
NM_003245	Homo sapiens transglutaminase 3 (E polypeptide, protein-glutamine-gamma-
NR 6 000250	glutamyltransferase) (TGM3), mRNA Homo sapiens transglutaminase 1 (K polypeptide epidermal type I, protein-
NM_000359	glutamine-gamma-glutamyltransferase) (TGM1), mRNA
NM 003243	Homo sapiens transforming growth factor, beta receptor III (betaglycan, 300kD)
14141_003243	(TGFBR3), mRNA
NM 003242	Homo sapiens transforming growth factor, beta receptor II (70-80kD)
14141_003242	(TGFBR2), mRNA
NM 000358	Homo sapiens transforming growth factor, beta-induced, 68kD (TGFBI), mRNA
NM 003239	Homo sapiens transforming growth factor, beta 3 (TGFB3), mRNA
NM 003239	Homo sapiens transforming growth factor, beta 2 (TGFB2), mRNA
NM 003236	Homo sapiens transforming growth factor, alpha (TGFA), mRNA
NM 003234	Homo sapiens transferrin receptor (p90, CD71) (TFRC), mRNA
NM 003227	Homo sapiens transferrin receptor 2 (TFR2), mRNA
NM 003226	Homo sapiens trefoil factor 3 (intestinal) (TFF3), mRNA
NM 003225	Homo sapiens trefoil factor 1 (breast cancer, estrogen-inducible sequence
1111_003223	expressed in) (TFF1), mRNA
NM_003224	Homo sapiens ADP-ribosylation factor related protein 1 (ARFRP1), mRNA
1111_003227	Table of the table to the table to the table to the table to the table to the table to the table to the table to the table to the table to the table to the table to the table to the table to the table to the table to the table to the table to table to the table to

NM_003219	Homo sapiens telomerase reverse transcriptase (TERT), mRNA
NM 003673	Homo sapiens titin-cap (telethonin) (TCAP), mRNA
NM 003217	Homo sapiens testis enhanced gene transcript (TEGT), mRNA
NM 003216	Homo sapiens thyrotrophic embryonic factor (TEF), mRNA
NM_003213	Homo sapiens TEA domain family member 4 (TEAD4), mRNA
NM 003211	Homo sapiens thymine-DNA glycosylase (TDG), mRNA
NM_003608	Homo sapiens G protein-coupled receptor 65 (GPR65), mRNA
NM 000355	Homo sapiens transcobalamin II; macrocytic anemia (TCN2), mRNA
NM_001062	Homo sapiens transcobalamin I (vitamin B12 binding protein, R binder family)
_	(TCN1), mRNA
NM_003202	Homo sapiens transcription factor 7 (T-cell specific, HMG-box) (TCF7), mRNA
NM_003201	Homo sapiens transcription factor 6-like 1 (mitochondrial transcription factor 1-
	like) (TCF6L1), mRNA
NM_003199	Homo sapiens transcription factor 4 (TCF4), mRNA
NM_003206	Homo sapiens transcription factor 21 (TCF21), mRNA
NM_000545	Homo sapiens transcription factor 1, hepatic; LF-B1, hepatic nuclear factor
	(HNF1), albumin proximal factor (TCF1), mRNA
NM_003198	Homo sapiens transcription elongation factor B (SIII), polypeptide 3 (110kD,
	elongin A) (TCEB3), mRNA
NM_001060	Homo sapiens thromboxane A2 receptor (TBXA2R), mRNA
NM_003194	Homo sapiens TATA box binding protein (TBP), mRNA
NM_003192	Homo sapiens tubulin-specific chaperone c (TBCC), mRNA
NM_000116	Homo sapiens tafazzin (cardiomyopathy, dilated 3A (X-linked); endocardial
	fibroelastosis 2; Barth syndrome) (TAZ), mRNA
NM_000353	Homo sapiens tyrosine aminotransferase (TAT), nuclear gene encoding
	mitochondrial protein, mRNA
NM_003191	Homo sapiens threonyl-tRNA synthetase (TARS), mRNA
NM_003190	Homo sapiens TAP binding protein (tapasin) (TAPBP), mRNA
NM_003189	Homo sapiens T-cell acute lymphocytic leukemia 1 (TAL1), mRNA
NM_003188	Homo sapiens mitogen-activated protein kinase kinase kinase 7 (MAP3K7),
377.6.000.407	mRNA
NM_003487	Homo sapiens TATA box binding protein (TBP)-associated factor, RNA
ND (002107	polymerase II, N, 68kD (RNA-binding protein 56) (TAF2N), mRNA
NM_003187	Homo sapiens TATA box binding protein (TBP)-associated factor, RNA polymerase II, G, 32kD (TAF2G), mRNA
NM 001057	Homo sapiens tachykinin receptor 2 (TACR2), mRNA
NM 003180	Homo sapiens synaptotagmin 5 (SYT5), mRNA
NM 003180	Homo sapiens synaptotaginin 3 (SY113), inkNA Homo sapiens synaptojanin 1 (SYNJ1), mRNA
NM 003490	Homo sapiens synapsin III (SYN3), mRNA
NM 003178	Homo sapiens synapsin II (SYN2), mRNA
NM 003178	Homo sapiens sylapsin it (STN2), indvA Homo sapiens spleen tyrosine kinase (SYK), mRNA
NM 003177	Homo sapiens synaptonemal complex protein 1 (SYCP1), mRNA
NM 003170	Homo sapiens surfeit 1 (SURF1), mRNA
NM 003172	Homo sapiens sulfett 1 (SORY 1), inclva. Homo sapiens sulfettasferase family, cytosolic, 2A, dehydroepiandrosterone
14141_003107	(DHEA) -preferring, member 1 (SULT2A1), mRNA
NM_001056	Homo sapiens sulfotransferase family, cytosolic, 1C, member 1 (SULT1C1),
14147_001020	mRNA
NM_001054	Homo sapiens sulfotransferase family, cytosolic, 1A, phenol-preferring, member
11117_001034	2 (SULT1A2), mRNA
NM 001055	Homo sapiens sulfotransferase family, cytosolic, 1A, phenol-preferring, member
11112_301033	1 (SULT1A1), mRNA
NM 003165	Homo sapiens syntaxin binding protein 1 (STXBP1), mRNA

NM_003163	Homo sapiens syntaxin 1B (STX1B), mRNA
NM_003159	Homo sapiens serine/threonine kinase 9 (STK9), mRNA
NM_003158	Homo sapiens serine/threonine kinase 6 (STK6), mRNA
NM_003157	Homo sapiens serine/threonine kinase 2 (STK2), mRNA
NM_003600	Homo sapiens serine/threonine kinase 15 (STK15), mRNA
NM_003160	Homo sapiens serine/threonine kinase 13 (aurora/IPL1-like) (STK13), mRNA
NM_003156	Homo sapiens stromal interaction molecule 1 (STIM1), mRNA
NM_003155	Homo sapiens stanniocalcin 1 (STC1), mRNA
NM_003877	Homo sapiens STAT induced STAT inhibitor-2 (STATI2), mRNA
NM_003154	Homo sapiens statherin (STATH), mRNA
NM_003153	Homo sapiens signal transducer and activator of transcription 6, interleukin-4 induced (STAT6), mRNA
NM_003152	Homo sapiens signal transducer and activator of transcription 5A (STAT5A), mRNA
NM_003151	Homo sapiens signal transducer and activator of transcription 4 (STAT4), mRNA
NM_003150	Homo sapiens signal transducer and activator of transcription 3 (acute-phase response factor) (STAT3), mRNA
NM 000349	Homo sapiens steroidogenic acute regulatory protein (STAR), mRNA
NM_003473	Homo sapiens signal transducing adaptor molecule (SH3 domain and ITAM motif) 1 (STAM), mRNA
NM_003149	Homo sapiens src homology three (SH3) and cysteine rich domain (STAC), mRNA
NM_001048	Homo sapiens somatostatin (SST), mRNA
NM 003146	Homo sapiens structure specific recognition protein 1 (SSRP1), mRNA
NM 003745	Homo sapiens JAK binding protein (SSI-1), mRNA
NM_001080	Homo sapiens aldehyde dehydrogenase 5 family, member A1 (succinate- semialdehyde dehydrogenase) (ALDH5A1), mRNA
NM_003139	Homo sapiens signal recognition particle receptor ('docking protein') (SRPR), mRNA
NM 003138	Homo sapiens SFRS protein kinase 2 (SRPK2), mRNA
NM 003135	Homo sapiens signal recognition particle 19kD (SRP19), mRNA
NM 003132	Homo sapiens spermidine synthase (SRM), mRNA
NM 003130	Homo sapiens sorcin (SRI), mRNA
NM_001047	Homo sapiens steroid-5-alpha-reductase, alpha polypeptide 1 (3-oxo-5 alpha-
	steroid delta 4-dehydrogenase alpha 1) (SRD5A1), mRNA
NM_003743	Homo sapiens nuclear receptor coactivator 1 (NCOA1), mRNA
NM_003128	Homo sapiens spectrin, beta, non-erythrocytic 1 (SPTBN1), mRNA
NM_003127	Homo sapiens spectrin, alpha, non-erythrocytic 1 (alpha-fodrin) (SPTAN1), mRNA
NM_003126	Homo sapiens spectrin, alpha, erythrocytic 1 (elliptocytosis 2) (SPTA1), mRNA
NM_003125	Homo sapiens small proline-rich protein 1B (cornifin) (SPRR1B), mRNA
NM_003124	Homo sapiens sepiapterin reductase (7,8-dihydrobiopterin:NADP+ oxidoreductase) (SPR), mRNA
NM_003123	Homo sapiens sialophorin (gpL115, leukosialin, CD43) (SPN), mRNA
NM_003121	Homo sapiens Spi-B transcription factor (Spi-1/PU.1 related) (SPIB), mRNA
NM_003120	Homo sapiens spleen focus forming virus (SFFV) proviral integration oncogene spi1 (SPI1), mRNA
NM_003119	Homo sapiens spastic paraplegia 7, paraplegin (pure and complicated autosomal recessive) (SPG7), mRNA
NM_003118	Homo sapiens secreted protein, acidic, cysteine-rich (osteonectin) (SPARC), mRNA
NM_003112	Homo sapiens Sp4 transcription factor (SP4), mRNA

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NM_003107	Homo sapiens SRY (sex determining region Y)-box 4 (SOX4), mRNA
	Homo sapiens SRY (sex determining region Y)-box 11 (SOX11), mRNA
	Homo sapiens sorbitol dehydrogenase (SORD), mRNA
	Homo sapiens superoxide dismutase 3, extracellular (SOD3), mRNA
NM_003794	Homo sapiens sorting nexin 4 (SNX4), mRNA
NM 003100	Homo sapiens sorting nexin 2 (SNX2), mRNA
NM 003094	Homo sapiens small nuclear ribonucleoprotein polypeptide E (SNRPE), mRNA
NM 003092	Homo sapiens small nuclear ribonucleoprotein polypeptide B" (SNRPB2),
-	mRNA
NM_003090	Homo sapiens small nuclear ribonucleoprotein polypeptide A' (SNRPA1),
-	mRNA
NM_003089	Homo sapiens small nuclear ribonucleoprotein 70kD polypeptide (RNP antigen)
-	(SNRP70), mRNA
NM 003498	Homo sapiens stannin (SNN), mRNA
NM 003087	Homo sapiens synuclein, gamma (breast cancer-specific protein 1) (SNCG),
	mRNA
NM 003083	Homo sapiens small nuclear RNA activating complex, polypeptide 2, 45kD
_	(SNAPC2), mRNA
NM 003082	Homo sapiens small nuclear RNA activating complex, polypeptide 1, 43kD
-	(SNAPC1), mRNA
NM 003081	Homo sapiens synaptosomal-associated protein, 25kD (SNAP25), mRNA
NM_003078	Homo sapiens SWI/SNF related, matrix associated, actin dependent regulator of
-	chromatin, subfamily d, member 3 (SMARCD3), mRNA
NM 003077	Homo sapiens SWI/SNF related, matrix associated, actin dependent regulator of
-	chromatin, subfamily d, member 2 (SMARCD2), mRNA
NM 003076	Homo sapiens SWI/SNF related, matrix associated, actin dependent regulator of
	chromatin, subfamily d, member 1 (SMARCD1), mRNA
NM 003075	Homo sapiens SWI/SNF related, matrix associated, actin dependent regulator of
-	chromatin, subfamily c, member 2 (SMARCC2), mRNA
NM_003074	Homo sapiens SWI/SNF related, matrix associated, actin dependent regulator of
	chromatin, subfamily c, member 1 (SMARCC1), mRNA
NM_003073	Homo sapiens SWI/SNF related, matrix associated, actin dependent regulator of
_	chromatin, subfamily b, member 1 (SMARCB1), mRNA
NM_003601	Homo sapiens SWI/SNF related, matrix associated, actin dependent regulator of
	chromatin, subfamily a, member 5 (SMARCA5), mRNA
NM_003071	Homo sapiens SWI/SNF related, matrix associated, actin dependent regulator of
_	chromatin, subfamily a, member 3 (SMARCA3), mRNA
NM_003070	Homo sapiens SWI/SNF related, matrix associated, actin dependent regulator of
	chromatin, subfamily a, member 2 (SMARCA2), mRNA
NM_003069	Homo sapiens SWI/SNF related, matrix associated, actin dependent regulator of
_	chromatin, subfamily a, member 1 (SMARCA1), mRNA
NM_003982	Homo sapiens solute carrier family 7 (cationic amino acid transporter, y+
-	system), member 7 (SLC7A7), mRNA
NM_003046	Homo sapiens solute carrier family 7 (cationic amino acid transporter, y+
	system), member 2 (SLC7A2), mRNA
NM_003045	Homo sapiens solute carrier family 7 (cationic amino acid transporter, y+
	system), member 1 (SLC7A1), mRNA
NM_003043	Homo sapiens solute carrier family 6 (neurotransmitter transporter, taurine),
	member 6 (SLC6A6), mRNA
NM_001045	Homo sapiens solute carrier family 6 (neurotransmitter transporter, serotonin),
	member 4 (SLC6A4), mRNA
NM_001044	Homo sapiens solute carrier family 6 (neurotransmitter transporter, dopamine),

	member 3 (SLC6A3), mRNA
NM 003042	Homo sapiens solute carrier family 6 (neurotransmitter transporter, GABA),
14141_003042	member 1 (SLC6A1), mRNA
ND4 002044	Homo sapiens solute carrier family 6 (neurotransmitter transporter,
NM_003044	betaine/GABA), member 12 (SLC6A12), mRNA
27.5.000452	Detaile/GADA), illefficer 12 (SECOA12), illeffix
NM_000453	Homo sapiens solute carrier family 5 (sodium iodide symporter), member 5
	(SLC5A5), mRNA
NM_003041	Homo sapiens solute carrier family 5 (sodium/glucose cotransporter), member 2
	(SLC5A2), mRNA
NM_000343	Homo sapiens solute carrier family 5 (sodium/glucose cotransporter), member 1
	(SLC5A1), mRNA
NM_003040	Homo sapiens solute carrier family 4, anion exchanger, member 2 (erythrocyte
_	membrane protein band 3-like 1) (SLC4A2), mRNA
NM 000342	Homo sapiens solute carrier family 4, anion exchanger, member 1 (erythrocyte
_	membrane protein band 3, Diego blood group) (SLC4A1), mRNA
NM_000341	Homo sapiens solute carrier family 3 (cystine, dibasic and neutral amino acid
_	transporters, activator of cystine, dibasic and neutral amino acid transport),
	member 1 (SLC3A1), mRNA
NM 001860	Homo sapiens solute carrier family 31 (copper transporters), member 2
11112_001000	(SLC31A2), mRNA
NM_001859	Homo sapiens solute carrier family 31 (copper transporters), member 1
14141_001033	(SLC31A1), mRNA
NB4 002020	Homo sapiens solute carrier family 2 (facilitated glucose transporter), member 5
NM_003039	
·	(SLC2A5), mRNA
NM_001042	Homo sapiens solute carrier family 2 (facilitated glucose transporter), member 4
	(SLC2A4), mRNA
NM_003705	Homo sapiens solute carrier family 25 (mitochondrial carrier, Aralar), member
	12 (SLC25A12), mRNA
NM_003060	Homo sapiens solute carrier family 22 (organic cation transporter), member 5
	(SLC22A5), mRNA
NM_003058	Homo sapiens solute carrier family 22 (organic cation transporter), member 2
	(SLC22A2), mRNA
NM_003057	Homo sapiens solute carrier family 22 (organic cation transporter), member 1
	(SLC22A1), mRNA
NM_003562	Homo sapiens solute carrier family 25 (mitochondrial carrier; oxoglutarate
_	carrier), member 11 (SLC25A11), mRNA
NM_003038	Homo sapiens solute carrier family 1 (glutamate/neutral amino acid transporter),
_	member 4 (SLC1A4), mRNA
NM 003056	Homo sapiens solute carrier family 19 (folate transporter), member 1
	(SLC19A1), mRNA
NM 003055	Homo sapiens solute carrier family 18 (vesicular acetylcholine), member 3
1111_005000	(SLC18A3), mRNA
NM 003054	Homo sapiens solute carrier family 18 (vesicular monoamine), member 2
14M1_003034	(SLC18A2), mRNA
NM_003053	Homo sapiens solute carrier family 18 (vesicular monoamine), member 1
14141 002022	•
ND (002052	(SLC18A1), mRNA
NM_003052	Homo sapiens solute carrier family 34 (sodium phosphate), member 1
	(SLC34A1), mRNA
NM_003051	Homo sapiens solute carrier family 16 (monocarboxylic acid transporters),
	member 1 (SLC16A1), mRNA
NM_003984	Homo sapiens solute carrier family 13 (sodium-dependent dicarboxylate
	transporter), member 2 (SLC13A2), mRNA

NM_000339	Homo sapiens solute carrier family 12 (sodium/chloride transporters), member 3 (SLC12A3), mRNA
NM_001046	Homo sapiens solute carrier family 12 (sodium/potassium/chloride transporters), member 2 (SLC12A2), mRNA
NM_000452	Homo sapiens solute carrier family 10 (sodium/bile acid cotransporter family), member 2 (SLC10A2), mRNA
NM_003049	Homo sapiens solute carrier family 10 (sodium/bile acid cotransporter family), member 1 (SLC10A1), mRNA
NM_003037	Homo sapiens signaling lymphocytic activation molecule (SLAM), mRNA
NM_003616	Homo sapiens survival of motor neuron protein interacting protein 1 (SIP1), mRNA
NM 003035	Homo sapiens TAL1 (SCL) interrupting locus (SIL), mRNA
NM_003032	Homo sapiens sialyltransferase 1 (beta-galactoside alpha-2,6-sialytransferase) (SIAT1), mRNA
NM 001041	Homo sapiens sucrase-isomaltase (SI), mRNA
NM 003027	Homo sapiens SH3-domain GRB2-like 3 (SH3GL3), mRNA
NM 003026	Homo sapiens SH3-domain GRB2-like 2 (SH3GL2), mRNA
NM 003025	Homo sapiens SH3-domain GRB2-like 1 (SH3GL1), mRNA
NM 003023	Homo sapiens SH3-domain binding protein 2 (SH3BP2), mRNA
NM_003022	Homo sapiens SH3 domain binding glutamic acid-rich protein like (SH3BGRL), mRNA
NM_000199	Homo sapiens N-sulfoglucosamine sulfohydrolase (sulfamidase) (SGSH), mRNA
NM_003020	Homo sapiens secretory granule, neuroendocrine protein 1 (7B2 protein) (SGNE1), mRNA
NM_000337	Homo sapiens sarcoglycan, delta (35kD dystrophin-associated glycoprotein) (SGCD), mRNA
NM_000232	Homo sapiens sarcoglycan, beta (43kD dystrophin-associated glycoprotein) (SGCB), mRNA
NM 003019	Homo sapiens surfactant, pulmonary-associated protein D (SFTPD), mRNA
NM 003018	Homo sapiens surfactant, pulmonary-associated protein C (SFTPC), mRNA
NM 000542	Homo sapiens surfactant, pulmonary-associated protein B (SFTPB), mRNA
NM 003011	Homo sapiens SET translocation (myeloid leukemia-associated) (SET), mRNA
NM 003010	Homo sapiens mitogen-activated protein kinase kinase 4 (MAP2K4), mRNA
NM 003009	Homo sapiens selenoprotein W, 1 (SEPW1), mRNA
NM 003008	Homo sapiens semenogelin II (SEMG2), mRNA
NM 003007	Homo sapiens semenogelin I (SEMG1), mRNA
NM_003966	Homo sapiens sema domain, seven thrombospondin repeats (type 1 and type 1-like), transmembrane domain (TM) and short cytoplasmic domain, (semaphorin) 5A (SEMA5A), mRNA
NM_003002	Homo sapiens succinate dehydrogenase complex, subunit D, integral membrane protein (SDHD), nuclear gene encoding mitochondrial protein, mRNA
NM_002999	Homo sapiens syndecan 4 (amphiglycan, ryudocan) (SDC4), mRNA
NM_002997	Homo sapiens syndecan 1 (SDC1), mRNA
NM_002996	Homo sapiens small inducible cytokine subfamily D (Cys-X3-Cys), member 1 (fractalkine, neurotactin) (SCYD1), mRNA
NM_003175	Homo sapiens small inducible cytokine subfamily C, member 2 (SCYC2), mRNA
NM_002993	Homo sapiens small inducible cytokine subfamily B (Cys-X-Cys), member 6 (granulocyte chemotactic protein 2) (SCYB6), mRNA
NM_002994	Homo sapiens small inducible cytokine subfamily B (Cys-X-Cys), member 5 (epithelial-derived neutrophil-activating peptide 78) (SCYB5), mRNA

NM 002985	Homo sapiens small inducible cytokine A5 (RANTES) (SCYA5), mRNA
NM 002991	Homo sapiens small inducible cytokine subfamily A (Cys-Cys), member 24
_	(SCYA24), mRNA
NM_002990	Homo sapiens small inducible cytokine subfamily A (Cys-Cys), member 22
	(SCYA22), mRNA
NM_002989	Homo sapiens small inducible cytokine subfamily A (Cys-Cys), member 21
11112_00_0	(SCYA21), mRNA
NM 002988	Homo sapiens small inducible cytokine subfamily A (Cys-Cys), member 18,
112.1_002.00	pulmonary and activation-regulated (SCYA18), mRNA
NM_002987	Homo sapiens small inducible cytokine subfamily A (Cys-Cys), member 17
- 12.12_00_0	(SCYA17), mRNA
NM 002986	Homo sapiens small inducible cytokine subfamily A (Cys-Cys), member 11
- ····-	(eotaxin) (SCYA11), mRNA
NM 002979	Homo sapiens sterol carrier protein 2 (SCP2), mRNA
NM 001039	Homo sapiens sodium channel, nonvoltage-gated 1, gamma (SCNN1G), mRNA
NM 002978	Homo sapiens sodium channel, nonvoltage-gated 1, delta (SCNN1D), mRNA
NM 001038	Homo sapiens sodium channel, nonvoltage-gated 1 alpha (SCNN1A), mRNA
NM_002977	Homo sapiens sodium channel, voltage-gated, type IX, alpha polypeptide
	(SCN9A), mRNA
NM 002976	Homo sapiens sodium channel, voltage-gated, type VI, alpha polypeptide
	(SCN6A), mRNA
NM_000334	Homo sapiens sodium channel, voltage-gated, type IV, alpha polypeptide
	(SCN4A), mRNA
NM 001037	Homo sapiens sodium channel, voltage-gated, type I, beta polypeptide (SCN1B),
	mRNA
NM 002975	Homo sapiens stem cell growth factor; lymphocyte secreted C-type lectin
	(SCGF), mRNA
NM 003843	Homo sapiens sciellin (SCEL), mRNA
NM 002973	Homo sapiens spinocerebellar ataxia 2 (olivopontocerebellar ataxia 2, autosomal
_	dominant, ataxin 2) (SCA2), mRNA
NM_000332	Homo sapiens spinocerebellar ataxia 1 (olivopontocerebellar ataxia 1, autosomal
_	dominant, ataxin 1) (SCA1), mRNA
NM_002971	Homo sapiens special AT-rich sequence binding protein 1 (binds to nuclear
_	matrix/scaffold-associating DNA's) (SATB1), mRNA
NM_002970	Homo sapiens spermidine/spermine N1-acetyltransferase (SAT), mRNA
NM_003870	Homo sapiens IQ motif containing GTPase activating protein 1 (IQGAP1),
.=	mRNA
NM_002967	Homo sapiens scaffold attachment factor B (SAFB), mRNA
NM_000331	Homo sapiens serum amyloid A1 (SAA1), mRNA
NM_001036	Homo sapiens ryanodine receptor 3 (RYR3), mRNA
NM_001035	Homo sapiens ryanodine receptor 2 (cardiac) (RYR2), mRNA
NM_002956	Homo sapiens restin (Reed-Steinberg cell-expressed intermediate filament-
	associated protein) (RSN), mRNA
NM_001033	Homo sapiens ribonucleotide reductase M1 polypeptide (RRM1), mRNA
NM_002955	Homo sapiens ras responsive element binding protein 1 (RREB1), mRNA
NM_003942	Homo sapiens ribosomal protein S6 kinase, 90kD, polypeptide 4 (RPS6KA4),
l	mRNA
NM_002953	Homo sapiens ribosomal protein S6 kinase, 90kD, polypeptide 1 (RPS6KA1),
	mRNA
NM_002951	Homo sapiens ribophorin II (RPN2), mRNA
NM_002950	Homo sapiens ribophorin I (RPN1), mRNA
NM_000329	Homo sapiens retinal pigment epithelium-specific protein (65kD) (RPE65),

NM_002947 NM_002946	mRNA Homo sapiens replication protein A3 (14kD) (RPA3), mRNA
NM_002946	
	The same a manufacture of the same of the
	Homo sapiens replication protein A2 (32kD) (RPA2), mRNA
	Homo sapiens replication protein A1 (70kD) (RPA1), mRNA
	Homo sapiens retinitis pigmentosa GTPase regulator (RPGR), mRNA
	Homo sapiens RAR-related orphan receptor A (RORA), mRNA
	Homo sapiens retinal outer segment membrane protein 1 (ROM1), mRNA
	Homo sapiens RNA (guanine-7-) methyltransferase (RNMT), mRNA
NM_002939	Homo sapiens ribonuclease/angiogenin inhibitor (RNH), mRNA
	Homo sapiens RNA guanylyltransferase and 5'-phosphatase (RNGTT), mRNA
NM_002938	Homo sapiens ring finger protein 4 (RNF4), mRNA
_	Homo sapiens ATP-binding cassette, sub-family E (OABP), member 1 (ABCE1), mRNA
	Homo sapiens ribonuclease H1 (RNASEH1), mRNA
-	Homo sapiens ribonuclease, RNase A family, 3 (eosinophil cationic protein) (RNASE3), mRNA
NM 002934	Homo sapiens ribonuclease, RNase A family, 2 (liver, eosinophil-derived
- ·	neurotoxin) (RNASE2), mRNA
	Homo sapiens RPB5-mediating protein (RMP), mRNA
NM 003821	Homo sapiens receptor-interacting serine-threonine kinase 2 (RIPK2), mRNA
	Homo sapiens LIM domain protein (RIL), mRNA
	Homo sapiens rhodopsin kinase (RHOK), mRNA
	Homo sapiens Rhesus blood group-associated glycoprotein (RHAG), mRNA
	Homo sapiens regulator of G-protein signalling 9 (RGS9), mRNA
	Homo sapiens regulator of G-protein signalling 5 (RGS5), mRNA
NM 002923	Homo sapiens regulator of G-protein signalling 2, 24kD (RGS2), mRNA
	Homo sapiens regulator of G-protein signalling 1 (RGS1), mRNA
NM 002928	Homo sapiens regulator of G-protein signalling 16 (RGS16), mRNA
NM 002926	Homo sapiens regulator of G-protein signalling 12 (RGS12), mRNA
NM 003834	Homo sapiens regulator of G-protein signalling 11 (RGS11), mRNA
NM 002921	Homo sapiens retinal G protein coupled receptor (RGR), mRNA
NM_000538	Homo sapiens regulatory factor X-associated protein (RFXAP), mRNA
NM_003721	Homo sapiens regulatory factor X-associated ankyrin-containing protein (RFXANK), mRNA
NM_002918	Homo sapiens regulatory factor X, 1 (influences HLA class II expression) (RFX1), mRNA
NM_002916	Homo sapiens replication factor C (activator 1) 4 (37kD) (RFC4), mRNA
NM_002915	Homo sapiens replication factor C (activator 1) 3 (38kD) (RFC3), mRNA
NM 002914	Homo sapiens replication factor C (activator 1) 2 (40kD) (RFC2), mRNA
NM_003704	Homo sapiens gene with multiple splice variants near HD locus on 4p16.3 (RES4-22), mRNA
NM_002908	Homo sapiens v-rel avian reticuloendotheliosis viral oncogene homolog (REL), mRNA
NM_002909	Homo sapiens regenerating islet-derived 1 alpha (pancreatic stone protein, pancreatic thread protein) (REG1A), mRNA
NM 000322	Homo sapiens retinal degeneration, slow (retinitis pigmentosa 7) (RDS), mRNA
NM 002905	Homo sapiens retinol dehydrogenase 5 (11-cisand 9-cis) (RDH5), mRNA
NM 002903	Homo sapiens recoverin (RCV1), mRNA
NM 002902	Homo sapiens reticulocalbin 2, EF-hand calcium binding domain (RCN2),
1111_00200	mRNA (1001001101112, E1 - Maile outstain onding domain (10012),
NM_002901	Homo sapiens reticulocalbin 1, EF-hand calcium binding domain (RCN1), mRNA

NM_002896	Homo sapiens RNA binding motif protein 4 (RBM4), mRNA
NM_002895	Homo sapiens retinoblastoma-like 1 (p107) (RBL1), mRNA
NM 000321	Homo sapiens retinoblastoma 1 (including osteosarcoma) (RB1), mRNA
NM 000966	Homo sapiens retinoic acid receptor, gamma (RARG), mRNA
NM 000964	Homo sapiens retinoic acid receptor, alpha (RARA), mRNA
NM 002885	Homo sapiens RAP1, GTPase activating protein 1 (RAP1GA1), mRNA
NM 002884	Homo sapiens RAP1A, member of RAS oncogene family (RAP1A), mRNA
NM 002883	Homo sapiens Ran GTPase activating protein 1 (RANGAP1), mRNA
NM_002881	Homo sapiens v-ral simian leukemia viral oncogene homolog B (ras related; GTP binding protein) (RALB), mRNA
NM_002871	Homo sapiens RAB interacting factor (RABIF), mRNA
NM 003929	Homo sapiens RAB7, member RAS oncogene family-like 1 (RAB7L1), mRNA
NM 002869	Homo sapiens RAB6, member RAS oncogene family (RAB6), mRNA
NM 002868	Homo sapiens RAB5B, member RAS oncogene family (RAB5B), mRNA
NM 002867	Homo sapiens RAB3B, member RAS oncogene family (RAB3B), mRNA
NM 002866	Homo sapiens RAB3A, member RAS oncogene family (RAB3A), mRNA
NM 002870	Homo sapiens RAB13, member RAS oncogene family (RAB13), mRNA
NM 000320	Homo sapiens quinoid dihydropteridine reductase (QDPR), mRNA
	Homo sapiens pregnancy-zone protein (PZP), mRNA
NM_002864	Homo sapiens phosphorylase, glycogen; liver (Hers disease, glycogen storage
NM_002863	disease type VI) (PYGL), mRNA
NM_002862	Homo sapiens phosphorylase, glycogen; brain (PYGB), nuclear gene encoding
	mitochondrial protein, mRNA
NM_002860	Homo sapiens pyrroline-5-carboxylate synthetase (glutamate gamma-
	semialdehyde synthetase) (PYCS), mRNA
NM_000319	Homo sapiens peroxisome receptor 1 (PXR1), mRNA
NM_002859	Homo sapiens paxillin (PXN), mRNA
NM_002857	Homo sapiens peroxisomal farnesylated protein (PXF), mRNA
NM_002854	Homo sapiens parvalbumin (PVALB), mRNA
NM_002852	Homo sapiens pentaxin-related gene, rapidly induced by IL-1 beta (PTX3), mRNA
NM 000317	Homo sapiens 6-pyruvoyltetrahydropterin synthase (PTS), mRNA
NM_002851	Homo sapiens protein tyrosine phosphatase, receptor-type, Z polypeptide 1 (PTPRZ1), mRNA
NM 002850	Homo sapiens protein tyrosine phosphatase, receptor type, S (PTPRS), mRNA
NM 002846	Homo sapiens protein tyrosine phosphatase, receptor type, N (PTPRN), mRNA
NM 002845	Homo sapiens protein tyrosine phosphatase, receptor type, M (PTPRM), mRNA
NM_002844	Homo sapiens protein tyrosine phosphatase, receptor type, K (PTPRK), mRNA
NM 002843	Homo sapiens protein tyrosine phosphatase, receptor type, J (PTPRJ), mRNA
NM_002842	Homo sapiens protein tyrosine phosphatase, receptor type, H (PTPRH), mRNA
NM 002840	Homo sapiens protein tyrosine phosphatase, receptor type, F (PTPRF), mRNA
NM 002839	Homo sapiens protein tyrosine phosphatase, receptor type, D (PTPRD), mRNA
NM 002824	Homo sapiens parathymosin (PTMS), mRNA
NM 002823	Homo sapiens prothymosin, alpha (gene sequence 28) (PTMA), mRNA
NM_000316	Homo sapiens parathyroid hormone receptor 1 (PTHR1), mRNA
NM_002820	Homo sapiens parathyroid hormone-like hormone (PTHLH), mRNA
NM_000315	Homo sapiens parathyroid hormone (PTH), mRNA
NM 000960	Homo sapiens prostaglandin I2 (prostacyclin) receptor (IP) (PTGIR), mRNA
NM 000959	Homo sapiens prostaglandin F receptor (FP) (PTGFR), mRNA
NM_000958	Homo sapiens prostaglandin E receptor 4 (subtype EP4) (PTGER4), mRNA
NM 000957	Homo sapiens prostaglandin E receptor 3 (subtype EP3) (PTGER3), mRNA
NM 000955	Homo sapiens prostaglandin E receptor 1 (subtype EP1), 42kD (PTGER1),
14147 000322	1 Itomo suprems prostagianum E receptor 1 (subtype Er 1), 1212 (1 TOEACT),

	mRNA
NM 000954	Homo sapiens prostaglandin D2 synthase (21kD, brain) (PTGDS), mRNA
NM_000314	Homo sapiens phosphatase and tensin homolog (mutated in multiple advanced cancers 1) (PTEN), mRNA
NM 000952	Homo sapiens platelet-activating factor receptor (PTAFR), mRNA
NM_002818	Homo sapiens proteasome (prosome, macropain) activator subunit 2 (PA28 beta)
1111_002010	(PSME2), mRNA
NM_002811	Homo sapiens proteasome (prosome, macropain) 26S subunit, non-ATPase, 7 (Mov34 homolog) (PSMD7), mRNA
NM_002806	Homo sapiens proteasome (prosome, macropain) 26S subunit, ATPase, 6 (PSMC6), mRNA
NM_002805	Homo sapiens proteasome (prosome, macropain) 26S subunit, ATPase, 5 (PSMC5), mRNA
NM_002804	Homo sapiens proteasome (prosome, macropain) 26S subunit, ATPase, 3 (PSMC3), mRNA
NM_002803	Homo sapiens proteasome (prosome, macropain) 26S subunit, ATPase, 2 (PSMC2), mRNA
NM_002802	Homo sapiens proteasome (prosome, macropain) 26S subunit, ATPase, 1 (PSMC1), mRNA
NM_002800	Homo sapiens proteasome (prosome, macropain) subunit, beta type, 9 (large multifunctional protease 2) (PSMB9), mRNA
NM_002799	Homo sapiens proteasome (prosome, macropain) subunit, beta type, 7 (PSMB7), mRNA
NM_002797	Homo sapiens proteasome (prosome, macropain) subunit, beta type, 5 (PSMB5), mRNA
NM_002796	Homo sapiens proteasome (prosome, macropain) subunit, beta type, 4 (PSMB4), mRNA
NM_002795	Homo sapiens proteasome (prosome, macropain) subunit, beta type, 3 (PSMB3), mRNA
NM_002794	Homo sapiens proteasome (prosome, macropain) subunit, beta type, 2 (PSMB2), mRNA
NM_002793	Homo sapiens proteasome (prosome, macropain) subunit, beta type, 1 (PSMB1), mRNA
NM_002801	Homo sapiens proteasome (prosome, macropain) subunit, beta type, 10 (PSMB10), mRNA
NM_002790	Homo sapiens proteasome (prosome, macropain) subunit, alpha type, 5 (PSMA5), mRNA
NM_002788	Homo sapiens proteasome (prosome, macropain) subunit, alpha type, 3 (PSMA3), mRNA
NM_002786	Homo sapiens proteasome (prosome, macropain) subunit, alpha type, 1 (PSMA1), mRNA
NM_002783	Homo sapiens pregnancy specific beta-1-glycoprotein 7 (PSG7), mRNA
NM_002781	Homo sapiens pregnancy specific beta-1-glycoprotein 5 (PSG5), mRNA
NM 002780	Homo sapiens pregnancy specific beta-1-glycoprotein 4 (PSG4), mRNA
NM_002785	Homo sapiens pregnancy specific beta-1-glycoprotein 11 (Note redefinition of symbol) (PSG11), mRNA
NM_002784	Homo sapiens pregnancy specific beta-1-glycoprotein 9 (PSG9), mRNA
NM_002779	Homo sapiens pleckstrin and Sec7 domain protein (PSD), mRNA
NM_002776	Homo sapiens kallikrein 10 (KLK10), mRNA
NM_002774	Homo sapiens kallikrein 6 (neurosin, zyme) (KLK6), mRNA
NM_002773	Homo sapiens protease, serine, 8 (prostasin) (PRSS8), mRNA
NM_002770	Homo sapiens protease, serine, 2 (trypsin 2) (PRSS2), mRNA

NM 002769	Homo sapiens protease, serine, 1 (trypsin 1) (PRSS1), mRNA
NM 003619	Homo sapiens protease, serine, 12 (neurotrypsin, motopsin) (PRSS12), mRNA
NM 002775	Homo sapiens protease, serine, 11 (IGF binding) (PRSS11), mRNA
NM 002767	Homo sapiens phosphoribosyl pyrophosphate synthetase-associated protein 2
_	(PRPSAP2), mRNA
NM 002766	Homo sapiens phosphoribosyl pyrophosphate synthetase-associated protein 1
_	(PRPSAP1), mRNA
NM 002765	Homo sapiens phosphoribosyl pyrophosphate synthetase 2 (PRPS2), mRNA
NM 002764	Homo sapiens phosphoribosyl pyrophosphate synthetase 1 (PRPS1), mRNA
NM 003891	Homo sapiens protein Z, vitamin K-dependent plasma glycoprotein (PROZ),
-	mRNA
NM 002763	Homo sapiens prospero-related homeobox 1 (PROX1), mRNA
NM_000313	Homo sapiens protein S (alpha) (PROS1), mRNA
NM_000312	Homo sapiens protein C (inactivator of coagulation factors Va and VIIIa)
	(PROC), mRNA
NM_002762	Homo sapiens protamine 2 (PRM2), mRNA
NM_002761	Homo sapiens protamine 1 (PRM1), mRNA
NM_000949	Homo sapiens prolactin receptor (PRLR), mRNA
NM_000948	Homo sapiens prolactin (PRL), mRNA
NM_002759	Homo sapiens protein kinase, interferon-inducible double stranded RNA
	dependent (PRKR), mRNA
NM_002756	Homo sapiens mitogen-activated protein kinase kinase 3 (MAP2K3), mRNA
NM_002749	Homo sapiens mitogen-activated protein kinase 7 (MAPK7), mRNA
NM_002745	Homo sapiens mitogen-activated protein kinase 1 (MAPK1), mRNA
NM_002751	Homo sapiens mitogen-activated protein kinase 11 (MAPK11), mRNA
NM_002753	Homo sapiens mitogen-activated protein kinase 10 (MAPK10), mRNA
NM_002743	Homo sapiens protein kinase C substrate 80K-H (PRKCSH), mRNA
NM_002742	Homo sapiens protein kinase C, mu (PRKCM), mRNA
NM_002741	Homo sapiens protein kinase C-like 1 (PRKCL1), mRNA
NM_002740	Homo sapiens protein kinase C, iota (PRKCI), mRNA
NM_002738	Homo sapiens protein kinase C, beta 1 (PRKCB1), mRNA
NM_002737	Homo sapiens protein kinase C, alpha (PRKCA), mRNA
NM_002736	Homo sapiens protein kinase, cAMP-dependent, regulatory, type II, beta
	(PRKAR2B), mRNA
NM_002734	Homo sapiens protein kinase, cAMP-dependent, regulatory, type I, alpha (tissue
	specific extinguisher 1) (PRKAR1A), mRNA
NM_002733	Homo sapiens protein kinase, AMP-activated, gamma 1 non-catalytic subunit (PRKAG1), mRNA
NM_002731	Homo sapiens protein kinase, cAMP-dependent, catalytic, beta (PRKACB),
	mRNA
NM 002730	Homo sapiens protein kinase, cAMP-dependent, catalytic, alpha (PRKACA),
	mRNA
NM 000947	Homo sapiens primase, polypeptide 2A (58kD) (PRIM2A), mRNA
NM 000946	Homo sapiens primase, polypeptide 1 (49kD) (PRIM1), mRNA
NM 002728	Homo sapiens proteoglycan 2, bone marrow (natural killer cell activator,
-	eosinophil granule major basic protein) (PRG2), mRNA
NM_002727	Homo sapiens proteoglycan 1, secretory granule (PRG1), mRNA
NM_002726	Homo sapiens prolyl endopeptidase (PREP), mRNA
NM_002725	Homo sapiens proline arginine-rich end leucine-rich repeat protein (PRELP),
	mRNA
NM_002723	Homo sapiens proline-rich protein BstNI subfamily 4 (PRB4), mRNA
NM 002722	Homo sapiens pancreatic polypeptide (PPY), mRNA

NM_000310	Homo sapiens palmitoyl-protein thioesterase 1 (ceroid-lipofuscinosis, neuronal 1, infantile) (PPT1), mRNA
NM_002720	Homo sapiens protein phosphatase 4 (formerly X), catalytic subunit (PPP4C), mRNA
NM_002719	Homo sapiens protein phosphatase 2, regulatory subunit B (B56), gamma isoform (PPP2R5C), mRNA
NM_002715	Homo sapiens protein phosphatase 2 (formerly 2A), catalytic subunit, alpha isoform (PPP2CA), mRNA
NM_002713	Homo sapiens protein phosphatase 1, regulatory (inhibitor) subunit 8 (PPP1R8), mRNA
NM 002712	Homo sapiens protein phosphatase 1, regulatory subunit 7 (PPP1R7), mRNA
NM 002714	Homo sapiens protein phosphatase 1, regulatory subunit 10 (PPP1R10), mRNA
NM_002710	Homo sapiens protein phosphatase 1, catalytic subunit, gamma isoform (PPP1CC), mRNA
NM_002709	Homo sapiens protein phosphatase 1, catalytic subunit, beta isoform (PPP1CB), mRNA
NM_002708	Homo sapiens protein phosphatase 1, catalytic subunit, alpha isoform (PPP1CA), mRNA
NM 000309	Homo sapiens protoporphyrinogen oxidase (PPOX), mRNA
NM_002706	Homo sapiens protein phosphatase 1B (formerly 2C), magnesium-dependent, beta isoform (PPM1B), mRNA
NM 002705	Homo sapiens periplakin (PPL), mRNA
NM 000943	Homo sapiens peptidylprolyl isomerase C (cyclophilin C) (PPIC), mRNA
NM_000308	Homo sapiens protective protein for beta-galactosidase (galactosialidosis) (PPGB), mRNA
NM 002703	Homo sapiens phosphoribosyl pyrophosphate amidotransferase (PPAT), mRNA
NM 003712	Homo sapiens phosphatidic acid phosphatase type 2C (PPAP2C), mRNA
NM 003713	Homo sapiens phosphatidic acid phosphatase type 2B (PPAP2B), mRNA
NM 003711	Homo sapiens phosphatidic acid phosphatase type 2A (PPAP2A), mRNA
NM_002702	Homo sapiens POU domain, class 6, transcription factor 1 (POU6F1), mRNA
NM_002701	Homo sapiens POU domain, class 5, transcription factor 1 (POU5F1), mRNA
NM_002700	Homo sapiens POU domain, class 4, transcription factor 3 (POU4F3), mRNA
NM_000307	Homo sapiens POU domain, class 3, transcription factor 4 (POU3F4), mRNA
NM_002699	Homo sapiens POU domain, class 3, transcription factor 1 (POU3F1), mRNA
NM_002697	Homo sapiens POU domain, class 2, transcription factor 1 (POU2F1), mRNA
NM_000306	Homo sapiens POU domain, class 1, transcription factor 1 (Pit1, growth hormone factor 1) (POU1F1), mRNA
NM_000446	Homo sapiens paraoxonase 1 (PON1), mRNA
NM_000939	Homo sapiens proopiomelanocortin (adrenocorticotropin/ beta-lipotropin/ alpha-melanocyte stimulating hormone/ beta-melanocyte stimulating hormone/ beta-endorphin) (POMC), mRNA
NM_002693	Homo sapiens polymerase (DNA directed), gamma (POLG), nuclear gene encoding mitochondrial protein, mRNA
NM_002692	Homo sapiens polymerase (DNA directed), epsilon 2 (POLE2), mRNA
NM_002691	Homo sapiens polymerase (DNA directed), delta 1, catalytic subunit (125kD) (POLD1), mRNA
NM_002690	Homo sapiens polymerase (DNA directed), beta (POLB), mRNA
NM_003967	Homo sapiens putative neurotransmitter receptor (PNR), mRNA
NM_002686	Homo sapiens phenylethanolamine N-methyltransferase (PNMT), mRNA
NM_002677	Homo sapiens peripheral myelin protein 2 (PMP2), mRNA
NM_000304	Homo sapiens peripheral myelin protein 22 (PMP22), mRNA
NM 002676	Homo sapiens phosphomannomutase 1 (PMM1), mRNA

NM_002674	Homo sapiens pro-melanin-concentrating hormone (PMCH), mRNA
NM_002668	Homo sapiens proteolipid protein 2 (colonic epithelium-enriched) (PLP2),
	mRNA
NM_000935	Homo sapiens procollagen-lysine, 2-oxoglutarate 5-dioxygenase (lysine
	hydroxylase) 2 (PLOD2), mRNA
NM 002667	Homo sapiens phospholamban (PLN), mRNA
NM 002666	Homo sapiens perilipin (PLIN), mRNA
NM 002665	Homo sapiens plasminogen-like (PLGL), mRNA
NM 000301	Homo sapiens plasminogen (PLG), mRNA
NM 000445	Homo sapiens plectin 1, intermediate filament binding protein, 500kD (PLEC1),
11112_000115	mRNA
NM 002663	Homo sapiens phospholipase D2 (PLD2), mRNA
NM 002662	Homo sapiens phospholipase D1, phophatidylcholine-specific (PLD1), mRNA
NM 002661	Homo sapiens phospholipase C, gamma 2 (phosphatidylinositol-specific)
1111_002001	(PLCG2), mRNA
NM 002660	Homo sapiens phospholipase C, gamma 1 (formerly subtype 148) (PLCG1),
1117_002000	mRNA
NM 000933	Homo sapiens phospholipase C, beta 4 (PLCB4), mRNA
NM 002659	Homo sapiens plasminogen activator, urokinase receptor (PLAUR), mRNA
NM_002658	Homo sapiens plasminogen activator, urokinase (PLAU), mRNA
NM 002655	Homo sapiens pleiomorphic adenoma gene 1 (PLAG1), mRNA
NM_000929	Homo sapiens phospholipase A2, group V (PLA2G5), mRNA
NM_003706	Homo sapiens phospholipase A2, group IVC (cytosolic, calcium-independent)
MM_003700	(PLA2G4C), mRNA
NM 000300	Homo sapiens phospholipase A2, group IIA (platelets, synovial fluid)
MM_000300	(PLA2G2A), nuclear gene encoding mitochondrial protein, mRNA
NM 003561	Homo sapiens phospholipase A2, group X (PLA2G10), mRNA
NM 002654	Homo sapiens pyruvate kinase, muscle (PKM2), mRNA
NM 003691	Homo sapiens serine/threonine kinase 16 (STK16), mRNA
	Homo sapiens polycystic kidney disease 1 (autosomal dominant) (PKD1),
NM_000296	mRNA
NM_003607	Homo sapiens Ser-Thr protein kinase related to the myotonic dystrophy protein
1411_003007	kinase (PK428), mRNA
NM 003678	Homo sapiens gene from NF2/meningioma region of 22q12 (PK1.3), mRNA
NM 000325	Homo sapiens paired-like homeodomain transcription factor 2 (PITX2), mRNA
NM 002653	Homo sapiens paired-like homeodomain transcription factor 1 (PITX1), mRNA
	Homo sapiens prolactin-induced protein (PIP), mRNA
NM_002652	Homo sapiens phosphatidylinositol-4-phosphate 5-kinase, type I, beta
NM_003558	(PIP5K1B), mRNA
ND4 002557	Homo sapiens phosphatidylinositol-4-phosphate 5-kinase, type I, alpha
NM_003557	(PIP5K1A), mRNA
ND (002746	Homo sapiens dynein, cytoplasmic, light polypeptide (PIN), mRNA
NM_003746	
NM_002648	Homo sapiens pim-1 oncogene (PIM1), mRNA
NM_002651	Homo sapiens phosphatidylinositol 4-kinase, catalytic, beta polypeptide
)D ('000(42	(PIK4CB), mRNA
NM_002643	Homo sapiens phosphatidylinositol glycan, class F (PIGF), mRNA
NM_002642	Homo sapiens phosphatidylinositol glycan, class C (PIGC), mRNA
NM_002638_	Homo sapiens protease inhibitor 3, skin-derived (SKALP) (PI3), mRNA
NM 000294	Homo sapiens phosphorylase kinase, gamma 2 (testis) (PHKG2), mRNA
NM_000293	Homo sapiens phosphorylase kinase, beta (PHKB), mRNA
NM_000292	Homo sapiens phosphorylase kinase, alpha 2 (liver) (PHKA2), mRNA
NM_002637	Homo sapiens phosphorylase kinase, alpha 1 (muscle) (PHKA1), mRNA

NM_000926	Homo sapiens progesterone receptor (PGR), mRNA
NM 002633	Homo sapiens phosphoglucomutase 1 (PGM1), mRNA
NM 000291	Homo sapiens phosphoglycerate kinase 1 (PGK1), mRNA
NM 002632	Homo sapiens placental growth factor, vascular endothelial growth factor-related
	protein (PGF), mRNA
NM 002631	Homo sapiens phosphogluconate dehydrogenase (PGD), mRNA
NM 002630	Homo sapiens progastricsin (pepsinogen C) (PGC), mRNA
NM_000290	Homo sapiens phosphoglycerate mutase 2 (muscle) (PGAM2), mRNA
NM_002629	Homo sapiens phosphoglycerate mutase 1 (brain) (PGAM1), mRNA
NM_000289	Homo sapiens phosphofructokinase, muscle (PFKM), mRNA
NM_002626	Homo sapiens phosphofructokinase, liver (PFKL), mRNA
NM 002625	Homo sapiens 6-phosphofructo-2-kinase/fructose-2,6-biphosphatase 1
l	(PFKFB1), mRNA
NM_002621	Homo sapiens properdin P factor, complement (PFC), mRNA
NM_002620	Homo sapiens platelet factor 4 variant 1 (PF4V1), mRNA
NM_002619	Homo sapiens platelet factor 4 (PF4), mRNA
NM_000288	Homo sapiens peroxisomal biogenesis factor 7 (PEX7), mRNA
NM_000287	Homo sapiens peroxisomal biogenesis factor 6 (PEX6), mRNA
NM_003630	Homo sapiens peroxisomal biogenesis factor 3 (PEX3), mRNA
NM_000466	Homo sapiens peroxisome biogenesis factor 1 (PEX1), mRNA
NM_002618	Homo sapiens peroxisome biogenesis factor 13 (PEX13), mRNA
NM_000442	Homo sapiens platelet/endothelial cell adhesion molecule (CD31 antigen)
	(PECAM1), mRNA
NM_002614	Homo sapiens PDZ domain containing 1 (PDZK1), mRNA
NM_003477	Homo sapiens Pyruvate dehydrogenase complex, lipoyl-containing component
	X; E3-binding protein (PDX1), mRNA
NM_002613	Homo sapiens 3-phosphoinositide dependent protein kinase-1 (PDPK1), mRNA
NM_002612	Homo sapiens pyruvate dehydrogenase kinase, isoenzyme 4 (PDK4), mRNA
NM_000925	Homo sapiens pyruvate dehydrogenase (lipoamide) beta (PDHB), mRNA
NM_000284	Homo sapiens pyruvate dehydrogenase (lipoamide) alpha 1 (PDHA1), mRNA
NM_000924	Homo sapiens phosphodiesterase IB, calmodulin-dependent (PDE1B), mRNA
NM_002606	Homo sapiens phosphodiesterase 9A (PDE9A), mRNA
NM_002602	Homo sapiens phosphodiesterase 6G, cGMP-specific, rod, gamma (PDE6G),
ND (000001	mRNA
NM_002601	Homo sapiens phosphodiesterase 6D, cGMP-specific, rod, delta (PDE6D),
NR 000001	mRNA
NM_000921	Homo sapiens phosphodiesterase 3A, cGMP-inhibited (PDE3A), mRNA
NM_002598	Homo sapiens programmed cell death 2 (PDCD2), mRNA Homo sapiens proprotein convertase subtilisin/kexin type 2 (PCSK2), mRNA
NM_002594	Homo sapiens proliferating cell nuclear antigen (PCNA), mRNA
NM_002592 NM_002591	Homo sapiens phosphoenolpyruvate carboxykinase 1 (soluble) (PCK1), mRNA
	Homo sapiens pre-B-cell leukemia transcription factor 2 (PBX2), mRNA
NM_002586 NM_002585	Homo sapiens pre-B-cell leukemia transcription factor 1 (PBX1), mRNA
NM_002583	Homo sapiens PRKC, apoptosis, WT1, regulator (PAWR), mRNA
NM_002582	Homo sapiens poly(A)-specific ribonuclease (deadenylation nuclease) (PARN),
11111_002302	mRNA
NM_003631	Homo sapiens poly (ADP-ribose) glycohydrolase (PARG), mRNA
NM_002580	Homo sapiens pancreatitis-associated protein (PAP), mRNA
NM 000919	Homo sapiens peptidylglycine alpha-amidating monooxygenase (PAM), mRNA
NM_002578	Homo sapiens p21 (CDKN1A)-activated kinase 3 (PAK3), mRNA
NM_002574	Homo sapiens peroxiredoxin 1 (PRDX1), mRNA
NM_002573	Homo sapiens platelet-activating factor acetylhydrolase, isoform lb, gamma
14147 007212	1 10110 sapiens platelet-activating factor acciyinytholase, isoform to, gaining

	subunit (29kD) (PAFAH1B3), mRNA
NM_002572	Homo sapiens platelet-activating factor acetylhydrolase, isoform Ib, beta subunit (30kD) (PAFAH1B2), mRNA
NM_002571	Homo sapiens progestagen-associated endometrial prot in (placental protein 14, pregnancy-associated endometrial alpha-2-globulin, alpha uterine protein) (PAEP), mRNA
NM_002569	Homo sapiens paired basic amino acid cleaving enzyme (furin, membrane associated receptor protein) (PACE), mRNA
NM 002570	Homo sapiens paired basic amino acid cleaving system 4 (PACE4), mRNA
NM 003900	Homo sapiens sequestosome 1 (SQSTM1), mRNA
NM_000918	Homo sapiens procollagen-proline, 2-oxoglutarate 4-dioxygenase (proline 4-hydroxylase), beta polypeptide (protein disulfide isomerase; thyroid hormone binding protein p55) (P4HB), mRNA
NM_000917	Homo sapiens procollagen-proline, 2-oxoglutarate 4-dioxygenase (proline 4-hydroxylase), alpha polypeptide I (P4HA1), mRNA
NM_002565	Homo sapiens pyrimidinergic receptor P2Y, G-protein coupled, 4 (P2RY4), mRNA
NM 002564	Homo sapiens purinergic receptor P2Y, G-protein coupled, 2 (P2RY2), mRNA
NM_002566	Homo sapiens purinergic receptor P2Y, G-protein coupled, 11 (P2RY11), mRNA
NM_002562	Homo sapiens purinergic receptor P2X, ligand-gated ion channel, 7 (P2RX7), mRNA
NM_002561	Homo sapiens purinergic receptor P2X, ligand-gated ion channel, 5 (P2RX5), mRNA
NM_002560	Homo sapiens purinergic receptor P2X, ligand-gated ion channel, 4 (P2RX4), mRNA
NM_002559	Homo sapiens purinergic receptor P2X, ligand-gated ion channel, 3 (P2RX3), mRNA
NM_002556	Homo sapiens oxysterol binding protein (OSBP), mRNA
NM 000608	Homo sapiens orosomucoid 2 (ORM2), mRNA
NM_003696	Homo sapiens olfactory receptor, family 6, subfamily A, member 1 (OR6A1), mRNA
NM_002550	Homo sapiens olfactory receptor, family 3, subfamily A, member 1 (OR3A1), mRNA
NM_002548	Homo sapiens olfactory receptor, family 1, subfamily D, member 2 (OR1D2), mRNA
NM_000914	Homo sapiens opioid receptor, mu 1 (OPRM1), mRNA
NM_000912	Homo sapiens opioid receptor, kappa 1 (OPRK1), mRNA
NM_000911	Homo sapiens opioid receptor, delta 1 (OPRD1), mRNA
NM_002544	Homo sapiens oligodendrocyte myelin glycoprotein (OMG), mRNA
NM_002543	Homo sapiens oxidised low density lipoprotein (lectin-like) receptor 1 (OLR1), mRNA
NM_003485	Homo sapiens G protein-coupled receptor 68 (GPR68), mRNA
NM_002540	Homo sapiens outer dense fibre of sperm tails 2 (ODF2), mRNA
NM_002533	Homo sapiens nuclear VCP-like (NVL), mRNA
NM 002531	Homo sapiens neurotensin receptor 1 (high affinity) (NTSR1), mRNA
NM 002530	Homo sapiens neurotrophic tyrosine kinase, receptor, type 3 (NTRK3), mRNA
NM_002526	Homo sapiens 5' nucleotidase (CD73) (NT5), mRNA
NM_003580	Homo sapiens neutral sphingomyelinase (N-SMase) activation associated factor (NSMAF), mRNA
NM_003633	Homo sapiens ectodermal-neural cortex (with BTB-like domain) (ENC1), mRNA

NM_003872	Homo sapiens neuropilin 2 (NRP2), mRNA
NM_003873	Homo sapiens neuropilin 1 (NRP1), mRNA
NM_003489	Homo sapiens nuclear receptor interacting protein 1 (NRIP1), mRNA
NM_002525	Homo sapiens nardilysin (N-arginine dibasic convertase) (NRD1), mRNA
NM_000905	Homo sapiens neuropeptide Y (NPY), mRNA
NM_000910	Homo sapiens neuropeptide Y receptor Y2 (NPY2R), mRNA
NM_000909	Homo sapiens neuropeptide Y receptor Y1 (NPY1R), mRNA
NM_002522	Homo sapiens neuronal pentraxin I (NPTX1), mRNA
NM_000908	Homo sapiens natriuretic peptide receptor C/guanylate cyclase C (atrionatriuretic peptide receptor C) (NPR3), mRNA
NM_000906	Homo sapiens natriuretic peptide receptor A/guanylate cyclase A (atrionatriuretic peptide receptor A) (NPR1), mRNA
NM 002521	Homo sapiens natriuretic peptide precursor B (NPPB), mRNA
NM 002519	Homo sapiens nuclear protein, ataxia-telangiectasia locus (NPAT), mRNA
NM 002518	Homo sapiens neuronal PAS domain protein 2 (NPAS2), mRNA
NM 002517	Homo sapiens neuronal PAS domain protein 1 (NPAS1), mRNA
NM 002514	Homo sapiens nephroblastoma overexpressed gene (NOV), mRNA
NM 003787	Homo sapiens nucleolar protein 4 (NOL4), mRNA
NM_003946	Homo sapiens nucleolar protein 3 (apoptosis repressor with CARD domain) (NOL3), mRNA
NM_003551	Homo sapiens non-metastatic cells 5, protein expressed in (nucleoside-diphosphate kinase) (NME5), mRNA
NM 002513	Homo sapiens non-metastatic cells 3, protein expressed in (NME3), mRNA
NM_002512	Homo sapiens non-metastatic cells 2, protein (NM23B) expressed in (NME2), nuclear gene encoding mitochondrial protein, mRNA
NM 002511	Homo sapiens neuromedin B receptor (NMBR), mRNA
NM 002510	Homo sapiens glycoprotein (transmembrane) nmb (GPNMB), mRNA
NM_003954	Homo sapiens mitogen-activated protein kinase kinase kinase 14 (MAP3K14), mRNA
NM 002508	Homo sapiens nidogen (enactin) (NID), mRNA
NM_002507	Homo sapiens nerve growth factor receptor (TNFR superfamily, member 16) (NGFR), mRNA
NM 002506	Homo sapiens nerve growth factor, beta polypeptide (NGFB), mRNA
NM_002503	Homo sapiens nuclear factor of kappa light polypeptide gene enhancer in B-cells inhibitor, beta (NFKBIB), mRNA
NM_002502	Homo sapiens nuclear factor of kappa light polypeptide gene enhancer in B-cells 2 (p49/p100) (NFKB2), mRNA
NM_002501	Homo sapiens nuclear factor I/X (CCAAT-binding transcription factor) (NFIX), mRNA
NM 002500	Homo sapiens neurogenic differentiation 1 (NEUROD1), mRNA
NM 002497	Homo sapiens NIMA (never in mitosis gene a)-related kinase 2 (NEK2), mRNA
NM_002496	Homo sapiens NADH dehydrogenase (ubiquinone) Fe-S protein 8 (23kD) (NADH-coenzyme Q reductase) (NDUFS8), mRNA
NM_002495	Homo sapiens NADH dehydrogenase (ubiquinone) Fe-S protein 4 (18kD) (NADH-coenzyme Q reductase) (NDUFS4), mRNA
NM_002494	Homo sapiens NADH dehydrogenase (ubiquinone) 1, subcomplex unknown, 1 (6kD, KFYI) (NDUFC1), mRNA
NM_002490	Homo sapiens NADH dehydrogenase (ubiquinone) 1 alpha subcomplex, 6
NM_002488	(14kD, B14) (NDUFA6), mRNA Homo sapiens NADH dehydrogenase (ubiquinone) 1 alpha subcomplex, 2 (8kD, B8) (NDUFA2), mRNA
NM 003635	Homo sapiens N-deacetylase/N-sulfotransferase (heparan glucosaminyl) 2
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	ADORS DIV
37.5 001540	(NDST2), mRNA
NM_001543	Homo sapiens N-deacetylase/N-sulfotransferase (heparan glucosaminyl) 1
N 5 000501	(NDST1), mRNA
NM_003581	Homo sapiens NCK adaptor protein 2 (NCK2), mRNA
NM_002486	Homo sapiens nuclear cap binding protein subunit 1, 80kD (NCBP1), mRNA
NM_002483	Homo sapiens carcinoembryonic antigen-related cell adhesion molecule 6 (non-specific cross reacting antigen) (CEACAM6), mRNA
NM_000662	Homo sapiens N-acetyltransferase 1 (arylamine N-acetyltransferase) (NAT1), mRNA
NM_000263	Homo sapiens N-acetylglucosaminidase, alpha- (Sanfilippo disease IIIB) (NAGLU), mRNA
NM_003871	Homo sapiens myelin transcription factor 2 (MYT2), mRNA
NM_003803	Homo sapiens myomesin 1 (skelemin) (185kD) (MYOM1), mRNA
NM 002479	Homo sapiens myogenin (myogenic factor 4) (MYOG), mRNA
NM_002472	Homo sapiens myosin, heavy polypeptide 8, skeletal muscle, perinatal (MYH8), mRNA
NM 002469	Homo sapiens myogenic factor 6 (herculin) (MYF6), mRNA
NM_002468	Homo sapiens myeloid differentiation primary response gene (88) (MYD88), mRNA
NM 002460	Homo sapiens interferon regulatory factor 4 (IRF4), mRNA
NM 002457	Homo sapiens mucin 2, intestinal/tracheal (MUC2), mRNA
NM 002456	Homo sapiens mucin 1, transmembrane (MUC1), mRNA
NM_002455	Homo sapiens metaxin 1 (MTX1), mRNA
NM_002453	Homo sapiens mitochondrial translational initiation factor 2 (MTIF2), nuclear
	gene encoding mitochondrial protein, mRNA
NM_002452	Homo sapiens nudix (nucleoside diphosphate linked moiety X)-type motif 1 (NUDT1), mRNA
NM_002450	Homo sapiens metallothionein 1L (MT1L), mRNA
NM_002447	Homo sapiens macrophage stimulating 1 receptor (c-met-related tyrosine kinase) (MST1R), mRNA
NM_002446	Homo sapiens mitogen-activated protein kinase kinase kinase 10 (MAP3K10), mRNA
NM 002445	Homo sapiens macrophage scavenger receptor 1 (MSR1), mRNA
NM 002444	Homo sapiens moesin (MSN), mRNA
NM_003879	Homo sapiens CASP8 and FADD-like apoptosis regulator (CFLAR), mRNA
NM_000530	Homo sapiens myelin protein zero (Charcot-Marie-Tooth neuropathy 1B) (MPZ), mRNA
NM_002437	Homo sapiens MpV17 transgene, murine homolog, glomerulosclerosis (MPV17), mRNA
NM_001932	Homo sapiens membrane protein, palmitoylated 3 (MAGUK p55 subfamily member 3) (MPP3), mRNA
NM_002435	Homo sapiens mannose phosphate isomerase (MPI), mRNA
NM_002434	Homo sapiens N-methylpurine-DNA glycosylase (MPG), mRNA
NM_003829	Homo sapiens multiple PDZ domain protein (MPDZ), mRNA
NM 003824	Homo sapiens Fas (TNFRSF6)-associated via death domain (FADD), mRNA
NM 002432	Homo sapiens myeloid cell nuclear differentiation antigen (MNDA), mRNA
NM_002431	Homo sapiens menage a trois 1 (CAK assembly factor) (MNAT1), mRNA
NM_002430	Homo sapiens meningioma (disrupted in balanced translocation) 1 (MN1), mRNA
NM_000901	Homo sapiens nuclear receptor subfamily 3, group C, member 2 (NR3C2), mRNA
NM_003482	Homo sapiens myeloid/lymphoid or mixed-lineage leukemia 2 (MLL2), mRNA

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NM_002419	Homo sapiens mitogen-activated protein kinase kinase kinase 11 (MAP3K11), mRNA
NM_002417	Homo sapiens antigen identified by monoclonal antibody Ki-67 (MKI67), mRNA
NM 002416	Homo sapiens monokine induced by gamma interferon (MIG), mRNA
NM 002415	Homo sapiens macrophage migration inhibitory factor (glycosylation-inhibiting
_	factor) (MIF), mRNA
NM_002413	Homo sapiens microsomal glutathione S-transferase 2 (MGST2), mRNA
NM_000900	Homo sapiens matrix Gla protein (MGP), mRNA
NM_002412	Homo sapiens O-6-methylguanine-DNA methyltransferase (MGMT), mRNA
NM_002407	Homo sapiens mammaglobin 2 (MGB2), mRNA
NM_002411_	Homo sapiens mammaglobin 1 (MGB1), mRNA
NM_002397	Homo sapiens MADS box transcription enhancer factor 2, polypeptide C
	(myocyte enhancer factor 2C) (MEF2C), mRNA
NM_002391	Homo sapiens midkine (neurite growth-promoting factor 2) (MDK), mRNA
NM_002387	Homo sapiens mutated in colorectal cancers (MCC), mRNA
NM_000529	Homo sapiens melanocortin 2 receptor (adrenocorticotropic hormone) (MC2R), mRNA
NM_002386	Homo sapiens melanocortin 1 receptor (alpha melanocyte stimulating hormone receptor) (MC1R), mRNA
NM_002385	Homo sapiens myelin basic protein (MBP), mRNA
NM_002382	Homo sapiens MAX protein (MAX), mRNA
NM_002378	Homo sapiens megakaryocyte-associated tyrosine kinase (MATK), mRNA
NM_002376	Homo sapiens MAP/microtubule affinity-regulating kinase 3 (MARK3), mRNA
NM_000898	Homo sapiens monoamine oxidase B (MAOB), nuclear gene encoding mitochondrial protein, mRNA
NM 003480	Homo sapiens Microfibril-associated glycoprotein-2 (MAGP2), mRNA
NM 002367	Homo sapiens melanoma antigen, family B, 4 (MAGEB4), mRNA
NM_002365	Homo sapiens melanoma antigen, family B, 3 (MAGEB3), mRNA
NM_002364	Homo sapiens melanoma antigen, family B, 2 (MAGEB2), mRNA
NM_002363	Homo sapiens melanoma antigen, family B, 1 (MAGEB1), mRNA
NM_002362	Homo sapiens melanoma antigen, family A, 4 (MAGEA4), mRNA
NM_003682	Homo sapiens MAP-kinase activating death domain (MADD), mRNA
NM_002357	Homo sapiens MAX dimerization protein (MAD), mRNA
NM_002350	Homo sapiens v-yes-1 Yamaguchi sarcoma viral related oncogene homolog (LYN), mRNA
NM 002349	Homo sapiens lymphocyte antigen 75 (LY75), mRNA
NM_002347	Homo sapiens lymphocyte antigen 6 complex, locus H (LY6H), mRNA
NM_002346	Homo sapiens lymphocyte antigen 6 complex, locus E (LY6E), mRNA
NM 002345	Homo sapiens lumican (LUM), mRNA
NM 002344	Homo sapiens leukocyte tyrosine kinase (LTK), mRNA
NM 002343	Homo sapiens lactotransferrin (LTF), mRNA
NM 000897	Homo sapiens leukotriene C4 synthase (LTC4S), mRNA
NM_003573	Homo sapiens latent transforming growth factor beta binding protein 4 (LTBP4), mRNA
NM_000752	Homo sapiens leukotriene b4 receptor (chemokine receptor-like 1) (LTB4R), mRNA
NM 000895	Homo sapiens leukotriene A4 hydrolase (LTA4H), mRNA
NM_002340	Homo sapiens lanosterol synthase (2,3-oxidosqualene-lanosterol cyclase) (LSS), mRNA
NM 002338	Homo sapiens limbic system-associated membrane protein (LSAMP), mRNA
NM_002337	Homo sapiens low density lipoprotein-related protein-associated protein 1

, 	(11 0 models line recently against departing 1) (I DDAD1) mDNA
27.5.00006	(alpha-2-macroglobulin receptor-associated protein 1) (LRPAP1), mRNA
NM 002336	Homo sapiens low density lipoprotein receptor-related protein 6 (LRP6), mRNA
NM_002319	Homo sapiens leucine-rich neuronal protein (LRN), mRNA
NM_002317	Homo sapiens lysyl oxidase (LOX), mRNA
NM_002316	Homo sapiens LIM homeobox transcription factor 1, beta (LMX1B), mRNA
NM_002315	Homo sapiens LIM domain only 1 (rhombotin 1) (LMO1), mRNA
NM_002312	Homo sapiens ligase IV, DNA, ATP-dependent (LIG4), mRNA
NM_002306	Homo sapiens lectin, galactoside-binding, soluble, 3 (galectin 3) (LGALS3), mRNA
NM 002303	Homo sapiens leptin receptor (LEPR), mRNA
NM 002302	Homo sapiens leukocyte cell-derived chemotaxin 2 (LECT2), mRNA
NM 001290	Homo sapiens LIM domain binding 2 (LDB2), mRNA
NM 003893	Homo sapiens LIM domain binding 1 (LDB1), mRNA
NM 002299	Homo sapiens lactase (LCT), mRNA
NM 002297	Homo sapiens lipocalin 1 (protein migrating faster than albumin, tear
<u>-</u>	prealbumin) (LCN1), mRNA
NM_002296	Homo sapiens lamin B receptor (LBR), mRNA
NM_002291	Homo sapiens laminin, beta 1 (LAMB1), mRNA
NM_002289	Homo sapiens lactalbumin, alpha- (LALBA), mRNA
NM 002273	Homo sapiens keratin 8 (KRT8), mRNA
NM 002276	Homo sapiens keratin 19 (KRT19), mRNA
NM 002275	Homo sapiens keratin 15 (KRT15), mRNA
NM 002274	Homo sapiens keratin 13 (KRT13), mRNA
NM 002265	Homo sapiens karyopherin (importin) beta 1 (KPNB1), mRNA
NM 002267	Homo sapiens karyopherin alpha 3 (importin alpha 4) (KPNA3), mRNA
NM_002266	Homo sapiens karyopherin alpha 2 (RAG cohort 1, importin alpha 1) (KPNA2), mRNA
NM 000893	Homo sapiens kininogen (KNG), mRNA
NM_003679	Homo sapiens kynurenine 3-monooxygenase (kynurenine 3-hydroxylase) (KMO), mRNA
NM_002258	Homo sapiens killer cell lectin-like receptor subfamily B, member 1 (KLRB1), mRNA
NM 002257	Homo sapiens kallikrein 1, renal/pancreas/salivary (KLK1), mRNA
NM 002256	Homo sapiens KiSS-1 metastasis-suppressor (KISS1), mRNA
NM_002255	Homo sapiens killer cell immunoglobulin-like receptor, two domains, long cytoplasmic tail, 4 (KIR2DL4), mRNA
NM 002254	Homo sapiens kinesin family member 3C (KIF3C), mRNA
NM_003958	Homo sapiens kinesin family method: 3C (kin 3C), mkd 41 Homo sapiens ring finger protein (C3HC4 type) 8 (RNF8), mRNA
NM 003685	Homo sapiens KH-type splicing regulatory protein (FUSE binding protein 2)
	(KHSRP), mRNA
NM_002252	Homo sapiens potassium voltage-gated channel, delayed-rectifier, subfamily S, member 3 (KCNS3), mRNA
NM_002250	Homo sapiens potassium intermediate/small conductance calcium-activated channel, subfamily N, member 4 (KCNN4), mRNA
NM_002249	Homo sapiens potassium intermediate/small conductance calcium-activated channel, subfamily N, member 3 (KCNN3), mRNA
NM_002247	Homo sapiens potassium large conductance calcium-activated channel, subfamily M, alpha member 1 (KCNMA1), mRNA
NM_002244	Homo sapiens potassium inwardly-rectifying channel, subfamily J, inhibitor 1 (KCNJN1), mRNA
NM_002240	Homo sapiens potassium inwardly-rectifying channel, subfamily J, member 6 (KCNJ6), mRNA

NM_002239	Homo sapiens potassium inwardly-rectifying channel, subfamily J, member 3 (KCNJ3), mRNA
NM_000891	Homo sapiens potassium inwardly-rectifying channel, subfamily J, member 2 (KCNJ2), mRNA
NM_002241	Homo sapiens potassium inwardly-rectifying channel, subfamily J, member 10 (KCNJ10), mRNA
NM_002238	Homo sapiens potassium voltage-gated channel, subfamily H (eag-related), member 1 (KCNH1), mRNA
NM_002237	Homo sapiens potassium voltage-gated channel, subfamily G, member 1 (KCNG1), mRNA
NM_002236	Homo sapiens potassium voltage-gated channel, subfamily F, member 1 (KCNF1), mRNA
NM_003636	Homo sapiens potassium voltage-gated channel, shaker-related subfamily, beta member 2 (KCNAB2), mRNA
NM_003471	Homo sapiens potassium voltage-gated channel, shaker-related subfamily, beta member 1 (KCNAB1), mRNA
NM_002235	Homo sapiens potassium voltage-gated channel, shaker-related subfamily, member 6 (KCNA6), mRNA
NM_002234	Homo sapiens potassium voltage-gated channel, shaker-related subfamily, member 5 (KCNA5), mRNA
NM_002233	Homo sapiens potassium voltage-gated channel, shaker-related subfamily, member 4 (KCNA4), mRNA
NM_002232	Homo sapiens potassium voltage-gated channel, shaker-related subfamily, member 3 (KCNA3), mRNA
NM_002229	Homo sapiens jun B proto-oncogene (JUNB), mRNA
NM_003666	Homo sapiens basic leucine zipper nuclear factor 1 (JEM-1) (BLZF1), mRNA
NM_002227	Homo sapiens Janus kinase 1 (a protein tyrosine kinase) (JAK1), mRNA
NM_003024	Homo sapiens intersectin 1 (SH3 domain protein) (ITSN1), mRNA
NM_002224	Homo sapiens inositol 1,4,5-triphosphate receptor, type 3 (TTPR3), mRNA
NM_002223	Homo sapiens inositol 1,4,5-triphosphate receptor, type 2 (TTPR2), mRNA
NM_002221	Homo sapiens inositol 1,4,5-trisphosphate 3-kinase B (ITPKB), mRNA
NM_002220	Homo sapiens inositol 1,4,5-trisphosphate 3-kinase A (ITPKA), mRNA
NM_002219	Homo sapiens integral membrane protein 1 (TTM1), mRNA
NM_002218	Homo sapiens inter-alpha (globulin) inhibitor H4 (plasma Kallikrein-sensitive glycoprotein) (ITIH4), mRNA
NM_002216	Homo sapiens inter-alpha (globulin) inhibitor, H2 polypeptide (ITIH2), mRNA
NM_002215	Homo sapiens inter-alpha (globulin) inhibitor, H1 polypeptide (ITIH1), mRNA
NM_000889	Homo sapiens integrin, beta 7 (ITGB7), mRNA
NM_002212	Homo sapiens integrin beta 4 binding protein (ITGB4BP), mRNA
NM_000213	Homo sapiens integrin, beta 4 (ITGB4), mRNA
NM_002211	Homo sapiens integrin, beta 1 (fibronectin receptor, beta polypeptide, antigen CD29 includes MDF2, MSK12) (ITGB1), mRNA
NM_002210	Homo sapiens integrin, alpha V (vitronectin receptor, alpha polypeptide, antigen CD51) (ITGAV), mRNA
NM_002209	Homo sapiens integrin, alpha L (antigen CD11A (p180), lymphocyte function-associated antigen 1; alpha polypeptide) (ITGAL), mRNA
NM_002206	Homo sapiens integrin, alpha 7 (ITGA7), mRNA
NM_002205	Homo sapiens integrin, alpha 5 (fibronectin receptor, alpha polypeptide) (ITGA5), mRNA
NM 003749	Homo sapiens insulin receptor substrate 2 (IRS2), mRNA
NM 001571 NM 002198	Homo sapiens interferon regulatory factor 3 (IRF3), mRNA Homo sapiens interferon regulatory factor 1 (IRF1), mRNA

NR (000106	Homo sapiens insulinoma-associated 1 (INSM1), mRNA
NM_002196	Homo sapiens insulin-like 4 (placenta) (INSL4), mRNA
NM_002195	Homo sapiens small inducible cytokine subfamily B (Cys-X-Cys), member 10
NM_001565	
37.6 000100	(SCYB10), mRNA Homo sapiens inhibin, beta A (activin A, activin AB alpha polypeptide)
NM_002192	
27.5.001564	(INHBA), mRNA Homo sapiens inhibitor of growth family, member 1-like (ING1L), mRNA
NM_001564	Homo sapiens inactivation escape 1 (INE1), mRNA
NM 003669	Homo sapiens IMP (inosine monophosphate) dehydrogenase 2 (IMPDH2),
NM_000884	mRNA
NM_000883	Homo sapiens IMP (inosine monophosphate) dehydrogenase 1 (IMPDH1),
1111_000005	mRNA
NM 001557	Homo sapiens interleukin 8 receptor, beta (IL8RB), mRNA
NM_000634	Homo sapiens interleukin 8 receptor, alpha (IL8RA), mRNA
NM 002185	Homo sapiens interleukin 7 receptor (IL7R), mRNA
NM 000880	Homo sapiens interleukin 7 (IL7), mRNA
NM 002184	Homo sapiens interleukin 6 signal transducer (gp130, oncostatin M receptor)
1111_002107	(IL6ST), mRNA
NM 000565	Homo sapiens interleukin 6 receptor (IL6R), mRNA
NM 000879	Homo sapiens interleukin 5 (colony-stimulating factor, eosinophil) (IL5), mRNA
NM 000589	Homo sapiens interleukin 4 (IL4), mRNA
NM 000588	Homo sapiens interleukin 3 (colony-stimulating factor, multiple) (IL3), mRNA
NM 000878	Homo sapiens interleukin 2 receptor, beta (IL2RB), mRNA
NM 003854	Homo sapiens interleukin 1 receptor-like 2 (IL1RL2), mRNA
NM 002182	Homo sapiens interleukin 1 receptor accessory protein (IL1RAP), mRNA
NM 000877	Homo sapiens interleukin 1 receptor, type I (IL1R1), mRNA
NM 003853	Homo sapiens interleukin 18 receptor accessory protein (IL18RAP), mRNA
NM 003855	Homo sapiens interleukin 18 receptor 1 (IL18R1), mRNA
NM 001562	Homo sapiens interleukin 18 (interferon-gamma-inducing factor) (IL18), mRNA
NM 002190	Homo sapiens interleukin 17 (cytotoxic T-lymphocyte-associated serine esterase
	8) (IL17), mRNA
NM 002189	Homo sapiens interleukin 15 receptor, alpha (IL15RA), mRNA
NM 002188	Homo sapiens interleukin 13 (IL13), mRNA
NM_001559	Homo sapiens interleukin 12 receptor, beta 2 (IL12RB2), mRNA
NM_002187	Homo sapiens interleukin 12B (natural killer cell stimulatory factor 2, cytotoxic
	lymphocyte maturation factor 2, p40) (IL12B), mRNA
NM_000882	Homo sapiens interleukin 12A (natural killer cell stimulatory factor 1, cytotoxic
	lymphocyte maturation factor 1, p35) (IL12A), mRNA
NM_000628	Homo sapiens interleukin 10 receptor, beta (IL10RB), mRNA
NM_001558	Homo sapiens interleukin 10 receptor, alpha (IL10RA), mRNA
NM_003639	Homo sapiens inhibitor of kappa light polypeptide gene enhancer in B-cells,
NR 002640	kinase gamma (IKBKG), mRNA Homo sapiens inhibitor of kappa light polypeptide gene enhancer in B-cells,
NM_003640	kinase complex-associated protein (IKBKAP), mRNA
NTM 001542	Homo sapiens immunoglobulin superfamily, member 3 (IGSF3), mRNA
NM 001542	Homo sapiens immunoglobulin superfamily, member 1 (IGSF1), mRNA
NM_001555	Homo sapiens immunoglobulin superiamity, memoer 1 (103x 1), micra Homo sapiens immunoglobulin mu binding protein 2 (IGHMBP2), mRNA
NM_002180	Homo sapiens immunoglobulin mu binding protein 2 (16HNBF2), mkVA Homo sapiens insulin-like growth factor binding protein 7 (IGFBP7), mRNA
NM_001553	Homo sapiens insulin-like growth factor binding protein 7 (IGFBF7), filed A Homo sapiens insulin-like growth factor binding protein 3 (IGFBP3), mRNA
NM_000598	There against insulin-like growth factor binding protein 1 (ICEPD1) mDNA
NM_000596	Homo sapiens insulin-like growth factor binding protein 1 (IGFBP1), mRNA
NM_001554	Homo sapiens cysteine-rich, angiogenic inducer, 61 (CYR61), mRNA
NM_000876	Homo sapiens insulin-like growth factor 2 receptor (IGF2R), mRNA

NM_001550	Homo sapiens interferon-related developmental regulator 1 (IFRD1), mRNA
NM_002177	Homo sapiens interferon, omega 1 (IFNW1), mRNA
NM_002176	Homo sapiens interferon, beta 1, fibr blast (IFNB1), mRNA
NM_000874	Homo sapiens interferon (alpha, beta and omega) receptor 2 (IFNAR2), mRNA
NM_002170	Homo sapiens interferon, alpha 8 (IFNA8), mRNA
NM 002169	Homo sapiens interferon, alpha 5 (IFNA5), mRNA
NM 002175	Homo sapiens interferon, alpha 21 (IFNA21), mRNA
NM_002173	Homo sapiens interferon, alpha 16 (IFNA16), mRNA
NM_002172	Homo sapiens interferon, alpha 14 (IFNA14), mRNA
NM_002171	Homo sapiens interferon, alpha 10 (IFNA10), mRNA
NM_001549	Homo sapiens interferon-induced protein with tetratricopeptide repeats 4 (IFIT4), mRNA
NM_001548	Homo sapiens interferon-induced protein with tetratricopeptide repeats 1 (IFIT1), mRNA
NM_003641	Homo sapiens interferon induced transmembrane protein 1 (9-27) (IFITM1), mRNA
NM 000204	Homo sapiens I factor (complement) (IF), mRNA
NM 002168	Homo sapiens isocitrate dehydrogenase 2 (NADP+), mitochondrial (IDH2),
<u> </u>	nuclear gene encoding mitochondrial protein, mRNA
NM_001546	Homo sapiens inhibitor of DNA binding 4, dominant negative helix-loop-helix
	protein (ID4), mRNA
NM_002166	Homo sapiens inhibitor of DNA binding 2, dominant negative helix-loop-helix
	protein (ID2), mRNA
NM_002165	Homo sapiens inhibitor of DNA binding 1, dominant negative helix-loop-helix protein (ID1), mRNA
NM_002160	Homo sapiens hexabrachion (tenascin C, cytotactin) (HXB), mRNA
NM_000871	Homo sapiens 5-hydroxytryptamine (serotonin) receptor 6 (HTR6), mRNA
NM_000869	Homo sapiens 5-hydroxytryptamine (serotonin) receptor 3A (HTR3A), mRNA
NM_000868	Homo sapiens 5-hydroxytryptamine (serotonin) receptor 2C (HTR2C), mRNA
NM_000867	Homo sapiens 5-hydroxytryptamine (serotonin) receptor 2B (HTR2B), mRNA
NM_000865	Homo sapiens 5-hydroxytryptamine (serotonin) receptor 1E (HTR1E), mRNA
NM_000864	Homo sapiens 5-hydroxytryptamine (serotonin) receptor 1D (HTR1D), mRNA
NM_000863	Homo sapiens 5-hydroxytryptamine (serotonin) receptor 1B (HTR1B), mRNA
NM_000524	Homo sapiens 5-hydroxytryptamine (serotonin) receptor 1A (HTR1A), mRNA
NM_002159	Homo sapiens histatin 1 (HTN1), mRNA
NM_002158	Homo sapiens human T-cell leukemia virus enhancer factor (HTLF), mRNA
NM_001541	Homo sapiens heat shock 27kD protein 2 (HSPB2), mRNA
NM_002155	Homo sapiens heat shock 70kD protein 6 (HSP70B') (HSPA6), mRNA
NM_001539	Homo sapiens heat shock protein, DNAJ-like 2 (HSJ2), mRNA
NM_000198	Homo sapiens hydroxy-delta-5-steroid dehydrogenase, 3 beta- and steroid delta-
	isomerase 2 (HSD3B2), mRNA
NM_000862	Homo sapiens hydroxy-delta-5-steroid dehydrogenase, 3 beta- and steroid delta-
776	isomerase 1 (HSD3B1), mRNA
NM_000414	Homo sapiens hydroxysteroid (17-beta) dehydrogenase 4 (HSD17B4), mRNA
NM_002153	Homo sapiens hydroxysteroid (17-beta) dehydrogenase 2 (HSD17B2), mRNA
NM_000413	Homo sapiens hydroxysteroid (17-beta) dehydrogenase 1 (HSD17B1), mRNA
NM_000196	Homo sapiens hydroxysteroid (11-beta) dehydrogenase 2 (HSD11B2), mRNA
NM_002151	Homo sapiens hepsin (transmembrane protease, serine 1) (HPN), mRNA
NM_000860	Homo sapiens hydroxyprostaglandin dehydrogenase 15-(NAD) (HPGD), mRNA
NM_002150	Homo sapiens 4-hydroxyphenylpyruvate dioxygenase (HPD), mRNA
NM_002143	Homo sapiens hippocalcin (HPCA), mRNA
NM_002148	Homo sapiens homeo box D10 (HOXD10), mRNA

NM 002145 Homo sapiens homeo box B2 (HOXB3), mRNA		
NM 002144 Homo sapiens homeo box B2 (HOXB2), mRNA	NM_002147	Homo sapiens homeo box B5 (HOXB5), mRNA
NM 002141 Homo sapiens homeo box B1 (HOXB1), mRNA		
NM 002142 Homo sapiens homeo box A9 (HOXA9), mRNA NM 002141 Homo sapiens homeo box A4 (HOXA4), mRNA NM 002522 Homo sapiens RNA binding motif protein, X chromosome (RBMX), mRNA NM 002139 Homo sapiens RNA binding motif protein, X chromosome (RBMX), mRNA NM 002135 Homo sapiens hepatocyte nuclear factor 4, alpha (HNP4A), mRNA NM 002135 Homo sapiens hepatocyte nuclear factor 4, alpha (HNP4A), mRNA NM 002131 Homo sapiens heme oxygenase (decycling) 1 (HMOX1), mRNA NM 002131 Homo sapiens high-mobility group (nonhistone chromosomal) protein isoforms I and Y (HMGIY), mRNA NM 002130 Homo sapiens high-mobility group (nonhistone chromosomal) protein isoforms I (HMGCS1), mRNA NM 002128 Homo sapiens high-mobility group (nonhistone chromosomal) protein 1 (HMGI), mRNA NM 001102 Homo sapiens high-mobility group (nonhistone chromosomal) protein 1 (HMGI), mRNA NM 001210 Homo sapiens hepatic leukemia factor (HLF), mRNA NM 001531 Homo sapiens major histocompatibility complex, class I-like sequence (HLALS), mRNA NM 002127 Homo sapiens major histocompatibility complex, class I, G (HLA-G), mRNA NM 001103 Homo sapiens major histocompatibility complex, class I, DQ beta I (HLA- DQBI), mRNA NM 001530 Homo sapiens HLA-G histocompatibility complex, class I, DQ beta I (HLA- DQBI), mRNA NM 001530 Homo sapiens hypoxia-inducible factor 1, alpha subunit (basic helix-loop-helix transcription factor) (HIFI IA), mRNA NM 001531 Homo sapiens hemochromatosis (HFE), mRNA NM 001532 Homo sapiens hemochromatosis (HFE), mRNA NM 003642 Homo sapiens hemochromatosis (HFE), mRNA NM 003141 Homo sapiens hemochromatosis (HFE), mRNA NM 003641 Homo sapiens hemochromatosis (HFE), mRNA NM 003642 Homo sapiens hemochromatosis (HFE), mRNA NM 003644 Homo sapiens hemochromatosis (HFE), mRNA NM 003547 Homo sapiens hemochromatosis (HFE), mRNA NM 003548 Homo sapiens histoine decarboxylase (HDC), mRNA NM 003549 Homo sapiens halistone family, member I (H4FI), mRNA NM 003540 Homo sapiens H4 histone family, member I (H4FI), mRNA NM 003541 Homo sapiens H3 histone family, member I		
NM 0001214 Homo sapiens homeo box A4 (HOXA4), mRNA NM 000522 Homo sapiens homeo box A13 (HOXAA13), mRNA NM 002139 Homo sapiens RNA binding motif protein, X chromosome (RBMX), mRNA NM 002135 Homo sapiens hepatocyte nuclear factor 4, alpha (HNF4A), mRNA NM 002135 Homo sapiens hepatocyte nuclear factor 4, alpha (HNF4A), mRNA NM 002131 Homo sapiens hepatocyte nuclear factor 4, alpha (HNF4A), mRNA NM 002131 Homo sapiens heme oxygenase (decycling) 1 (HMOX1), mRNA NM 002131 Homo sapiens heme oxygenase (decycling) 1 (HMOX1), mRNA NM 002131 Homo sapiens high-mobility group (nonhistone chromosomal) protein isoforms I and Y (HMGIY), mRNA NM 002130 Homo sapiens high-mobility group (nonhistone chromosomal) protein 1 (HMGCS1), mRNA NM 002128 Homo sapiens high-mobility group (nonhistone chromosomal) protein 1 (HMGI), mRNA NM 00190 Homo sapiens high-mobility group (nonhistone chromosomal) protein 1 (HMGI), mRNA NM 00191 Homo sapiens high-mobility group (nonhistone chromosomal) protein 1 (HMGI), mRNA NM 001531 Homo sapiens hepatic leukemia factor (HLF), mRNA NM 001531 Homo sapiens hepatic leukemia factor (HLF), mRNA NM 002126 Homo sapiens hepatic leukemia factor (HLF), mRNA NM 002127 Homo sapiens HLA-G histocompatibility complex, class I, G (HLA-G), mRNA NM 002127 Homo sapiens HIA-G histocompatibility antigen, class I, G (HLA-G), mRNA NM 001530 Homo sapiens hypoxia-inducible factor 1, alpha subunit (basic helix-loop-helix transcription factor) (HIF1A), mRNA NM 001530 Homo sapiens homogentisate 1,2-dioxygenase (homogentisate oxidase) (HGD), mRNA NM 001531 Homo sapiens hemochromatosis (HFE), mRNA NM 00165 Homo sapiens hemochromatosis (HFE), mRNA NM 00164 Homo sapiens histone acetyltransferase 1 (HAT1), mRNA NM 003865 Homo sapiens histone acetyltransferase 1 (HAT1), mRNA NM 003151 Homo sapiens hydroxyacyl-Coenzyme A dehydrogenase/3-ketoacyl-Coenzyme A thiolase/enoyl-Coenzyme A hydratase (trifunctional protein), beta subunit (HADHB), mRNA NM 003544 Homo sapiens H4 histone family, member I (H4FI), mRNA NM 003547 Homo sapiens H3 hist		
NM 002129 Homo sapiens homeo box A13 (HOXA13), mRNA NM 002139 Homo sapiens RNA binding motif protein, X chromosome (RBMX), mRNA NM 000457 Homo sapiens nuclear receptor subfamily 4, group A, member 1 (NR4A1), mRNA NM_002135 Homo sapiens heme oxygenase (decycling) 1 (HMOX1), mRNA NM_002131 Homo sapiens heme oxygenase (decycling) 1 (HMOX1), mRNA NM_002131 Homo sapiens high-mobility group (nonhistone chromosomal) protein isoforms 1 and Y (HMGIY), mRNA NM_002130 Homo sapiens high-mobility group (nonhistone chromosomal) protein 1 (HMGS1), mRNA NM_002130 Homo sapiens high-mobility group (nonhistone chromosomal) protein 1 (HMGCS1), mRNA NM_001126 Homo sapiens high-mobility group (nonhistone chromosomal) protein 1 (HMGI), mRNA NM_00190 Homo sapiens hydroxymethylbilane synthase (HMBS), mRNA NM_00151 Homo sapiens hepatic leukemia factor (HLF), mRNA NM_001531 Homo sapiens hepatic leukemia factor (HLF), mRNA NM_001212 Homo sapiens major histocompatibility complex, class I, G (HLA-G), mRNA NM_001213 Homo sapiens major histocompatibility complex, class I, DQ beta 1 (HLA-DQB1), mRNA NM_001530 Homo sapiens major histocompatibility complex, class I, DQ beta 1 (HLA-DQB1), mRNA NM_001530 Homo sapiens hypoxia-inducible factor 1, alpha subunit (basic helix-loop-helix transcription factor) (HIF1A), mRNA NM_001530 Homo sapiens homogentisate 1,2-dioxygenase (homogentisate oxidase) (HGD), mRNA NM_001520 Homo sapiens hemochromatosis (HFE), mRNA NM_00166 Homo sapiens hemochromatosis (HFE), mRNA NM_00166 Homo sapiens hemochromatosis (HFE), mRNA NM_00170 Homo sapiens hemochromatosis (HFE), mRNA NM_001531 Homo sapiens histidine decarboxylase (HDC), mRNA NM_001540 Homo sapiens hemochromatosis (HFE), mRNA NM_001541 Homo sapiens hemochromatosis (HFE), mRNA NM_001541 Homo sapiens histidine decarboxylase (HGC), mRNA NM_001541 Homo sapiens histidine decarboxylase (HGC), mRNA NM_003544 Homo sapiens hyaluronan synthase 1 (HAS1), mRNA NM_003545 Homo sapiens Halistone family, member I (H4FI), mRNA NM_003546 Homo sapiens Halistone family, member I (H4F	NM_002142	· · · · · · · · · · · · · · · · · · ·
NM 002139 Homo sapiens RNA binding motif protein, X chromosome (RBMX), mRNA NM 000457 Homo sapiens hepatocyte nuclear factor 4, alpha (HNR4A), mRNA NM_002135 Homo sapiens hepatocyte nuclear factor 4, alpha (HNR4A), mRNA NM_002131 Homo sapiens heme oxygenase (decycling) 1 (HMOX1), mRNA NM_002131 Homo sapiens heme oxygenase (decycling) 1 (HMOX1), mRNA NM_002131 Homo sapiens high-mobility group (nonhistone chromosomal) protein isoforms I and Y (HMGIY), mRNA NM_002130 Homo sapiens 3-hydroxy-3-methylglutaryl-Coenzyme A synthase 1 (soluble) (HMGCS1), mRNA NM_002128 Homo sapiens high-mobility group (nonhistone chromosomal) protein 1 (HMG1), mRNA NM_00190 Homo sapiens hydroxymethylbilane synthase (HMBS), mRNA NM_00150 Homo sapiens hydroxymethylbilane synthase (HMBS), mRNA NM_00151 Homo sapiens hydroxymethylbilane synthase (HMBS), mRNA NM_001521 Homo sapiens hydroxymethylbilane synthase (HMBS), mRNA NM_001531 Homo sapiens hydroxymethylbilane synthase (HMBS), mRNA NM_001532 Homo sapiens HLA-G histocompatibility complex, class I, G (HLA-G), mRNA NM_001530 Homo sapiens hydroxymethylbilane synthase (HMBS), mRNA NM_001530 Homo sapiens hydroxymethylbilane synthase (HMBS), mRNA NM_001530 Homo sapiens hydroxia-inducible factor 1, alpha subunit (basic helix-loop-helix transcription factor) (HIF1A), mRNA NM_001528 Homo sapiens HGF activator (HGFAC), mRNA NM_000187 Homo sapiens hemochromatosis (HFE), mRNA NM_000188 Homo sapiens Hfactor 1 (complement) (HF1), mRNA NM_000189 Homo sapiens hydroxymethylane (HGC), mRNA NM_000110 Homo sapiens hydroxymethylane (HGC), mRNA NM_000110 Homo sapiens hydroxymethylane (HGC), mRNA NM_000110 Homo sapiens hydroxymethylane (HGAS1), mRNA NM_000180 Homo sapiens hydroxymethylane (HGAS1), mRNA NM_000181 Homo sapiens hydroxymethylane (HGAS1), mRNA NM_000182 Homo sapiens hydroxymethylane (HGAS1), mRNA NM_000183 Homo sapiens hydroxymethylane (HGAS1), mRNA NM_000184 Homo sapiens hydroxymethylane (HGAS1), mRNA NM_000185 Homo sapiens H3 histone family, member I (H4FI), mRNA NM_00354 Homo sapiens H3 histone fa	NM_002141	<u></u>
NM 002135	NM_000522	
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NM 000410 Homo sapiens hemochromatosis (HFE), mRNA NM 000186 Homo sapiens H factor 1 (complement) (HF1), mRNA NM 003865 Homo sapiens homeo box (expressed in ES cells) 1 (HESX1), mRNA NM 002112 Homo sapiens histidine decarboxylase (HDC), mRNA NM 002110 Homo sapiens hemopoietic cell kinase (HCK), mRNA NM 003642 Homo sapiens histone acetyltransferase 1 (HAT1), mRNA NM 001523 Homo sapiens hydroxyacyl-Coenzyme A dehydrogenase/3-ketoacyl-Coenzyme A thiolase/enoyl-Coenzyme A hydratase (trifunctional protein), beta subunit (HADHB), mRNA NM 000182 Homo sapiens hydroxyacyl-Coenzyme A dehydrogenase/3-ketoacyl-Coenzyme A thiolase/enoyl-Coenzyme A hydratase (trifunctional protein), alpha subunit (HADHA), mRNA NM 003548 Homo sapiens H4 histone, family 2 (H4F2), mRNA NM 003547 Homo sapiens H4 histone family, member L (H4FL), mRNA NM 003544 Homo sapiens H4 histone family, member I (H4FI), mRNA NM 003537 Homo sapiens H3 histone family, member T (H3FT), mRNA NM 003534 Homo sapiens H3 histone family, member L (H3FL), mRNA NM 003534 Homo sapiens H3 histone family, member H (H3FH), mRNA NM 003534 Homo sapiens H3 histone family, member D (H3FD), mRNA NM 003531 Homo sapiens H3 histone family, member D (H3FD), mRNA NM 003531 Homo sapiens H3 histone family, member D (H3FD), mRNA		Homo sapiens homogentisate 1,2-dioxygenase (homogentisate oxidase) (HGD),
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NM 003642 Homo sapiens histone acetyltransferase 1 (HAT1), mRNA NM 001523 Homo sapiens hyaluronan synthase 1 (HAS1), mRNA NM 00183 Homo sapiens hydroxyacyl-Coenzyme A dehydrogenase/3-ketoacyl-Coenzyme A thiolase/enoyl-Coenzyme A hydratase (trifunctional protein), beta subunit (HADHB), mRNA NM 000182 Homo sapiens hydroxyacyl-Coenzyme A dehydrogenase/3-ketoacyl-Coenzyme A thiolase/enoyl-Coenzyme A hydratase (trifunctional protein), alpha subunit (HADHA), mRNA NM 003548 Homo sapiens H4 histone, family 2 (H4F2), mRNA NM 003547 Homo sapiens H4 histone family, member L (H4FL), mRNA NM 003544 Homo sapiens H4 histone family, member I (H4FI), mRNA NM 003545 Homo sapiens H3 histone family, member T (H3FT), mRNA NM 003546 Homo sapiens H3 histone family, member L (H3FL), mRNA NM 003537 Homo sapiens H3 histone family, member H (H3FH), mRNA NM 003534 Homo sapiens H3 histone family, member D (H3FD), mRNA NM 003531 Homo sapiens H3 histone family, member D (H3FD), mRNA NM 003531 Homo sapiens H3 histone family, member D (H3FD), mRNA		
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NM 003531 Homo sapiens H3 histone family, member C (H3FC), mRNA		
The state of the s	NM_003530	Homo sapiens H3 histone family, member B (H3FB), mRNA

	TYO I C. II A CICEAN DNA
NM_003529	Homo sapiens H3 histone family, member A (H3FA), mRNA
NM_002107	Homo sapiens H3 histone, family 3A (H3F3A), mRNA
NM_003528	Homo sapiens H2B histone family, member Q (H2BFQ), mRNA
NM_003526	Homo sapiens H2B histone family, member L (H2BFL), mRNA
NM_003525	Homo sapiens H2B histone family, member K (H2BFK), mRNA
NM 003524	Homo sapiens H2B histone family, member J (H2BFJ), mRNA
NM 003523	Homo sapiens H2B histone family, member H (H2BFH), mRNA
NM 003522	Homo sapiens H2B histone family, member G (H2BFG), mRNA
NM 003518	Homo sapiens H2B histone family, member A (H2BFA), mRNA
NM 002106	Homo sapiens H2A histone family, member Z (H2AFZ), mRNA
NM 003516	Homo sapiens H2A histone family, member O (H2AFO), mRNA
NM 003513	Homo sapiens H2A histone family, member M (H2AFM), mRNA
NM 003512	Homo sapiens H2A histone family, member L (H2AFL), mRNA
NM 003612	Homo sapiens sema domain, immunoglobulin domain (Ig), and GPI membrane
	anchor, (semaphorin) 7A (SEMA7A), mRNA
NM_002104	Homo sapiens granzyme K (serine protease, granzyme 3; tryptase II) (GZMK), mRNA
NM 002103	Homo sapiens glycogen synthase 1 (muscle) (GYS1), mRNA
NM 002102	Homo sapiens glycophorin E (GYPE), mRNA
NM 000181	Homo sapiens glucuronidase, beta (GUSB), mRNA
NM 000858	Homo sapiens guanylate kinase 1 (GUK1), mRNA
NM 001522	Homo sapiens guanylate cyclase 2F, retinal (GUCY2F), mRNA
NM 000180	Homo sapiens guanylate cyclase 2D, membrane (retina-specific) (GUCY2D),
	mRNA
NM_000857	Homo sapiens guanylate cyclase 1, soluble, beta 3 (GUCY1B3), mRNA
NM_000856	Homo sapiens guanylate cyclase 1, soluble, alpha 3 (GUCY1A3), mRNA
NM_000855	Homo sapiens guanylate cyclase 1, soluble, alpha 2 (GUCY1A2), mRNA
NM_000409	Homo sapiens guanylate cyclase activator 1A (retina) (GUCA1A), mRNA
NM_001517	Homo sapiens general transcription factor IIH, polypeptide 4 (52kD subunit) (GTF2H4), mRNA
NM_002096	Homo sapiens general transcription factor IIF, polypeptide 1 (74kD subunit) (GTF2F1), mRNA
NM_002095	Homo sapiens general transcription factor IIE, polypeptide 2 (beta subunit, 34kD) (GTF2E2), mRNA
NM_001513	Homo sapiens glutathione transferase zeta 1 (maleylacetoacetate isomerase) (GSTZ1), mRNA
NM 000853	Homo sapiens glutathione S-transferase theta 1 (GSTT1), mRNA
NM 000851	Homo sapiens glutathione S-transferase M5 (GSTM5), mRNA
NM 000850	Homo sapiens glutathione S-transferase M4 (GSTM4), mRNA
NM 000849	Homo sapiens glutathione S-transferase M3 (brain) (GSTM3), mRNA
NM 000848	Homo sapiens glutathione S-transferase M2 (muscle) (GSTM2), mRNA
NM 001512	Homo sapiens glutathione S-transferase A4 (GSTA4), mRNA
NM 000846	Homo sapiens glutathione S-transferase A2 (GSTA2), mRNA
NM 000178	Homo sapiens glutathione synthetase (GSS), mRNA
NM 002094	Homo sapiens G1 to S phase transition 1 (GSPT1), mRNA
NM 000177	Homo sapiens gelsolin (amyloidosis, Finnish type) (GSN), mRNA
NM 002093	Homo sapiens glycogen synthase kinase 3 beta (GSK3B), mRNA
NM_002093	Homo sapiens G-rich RNA sequence binding factor 1 (GRSF1), mRNA
	Homo sapiens gastrin-releasing peptide (GRP), mRNA
NM 002091	Homo sapiens GRO3 oncogene (GRO3), mRNA
NM 002090	Homo sapiens GRO2 oncogene (GRO2), mRNA
NM 002089	Homo sapiens GRO2 oncogene (GRO2), interval Homo sapiens GRO1 oncogene (melanoma growth stimulating activity, alpha)
NM_001511	nomo saprens OROI oncogene (meianoma growm summaring activity, aipha)

	(GRO1), mRNA
NM 002087	Homo sapiens granulin (GRN), mRNA
NM 000845	Homo sapiens glutamate receptor, metabotropic 8 (GRM8), mRNA
NM 000844	Homo sapiens glutamate receptor, metabotropic 7 (GRM7), mRNA
NM_000841	Homo sapiens glutamate receptor, metabotropic 4 (GRM4), mRNA
NM 000840	Homo sapiens glutamate receptor, metabotropic 3 (GRM3), mRNA
NM 000176	Homo sapiens nuclear receptor subfamily 3, group C, member 1 (NR3C1),
_ _	mRNA
NM_000831	Homo sapiens glutamate receptor, ionotropic, kainate 3 (GRIK3), mRNA
NM_000830	Homo sapiens glutamate receptor, ionotropic, kainate 1 (GRIK1), mRNA
NM_002086	Homo sapiens growth factor receptor-bound protein 2 (GRB2), mRNA
NM_002085	Homo sapiens glutathione peroxidase 4 (phospholipid hydroperoxidase) (GPX4), mRNA
NM 002083	Homo sapiens glutathione peroxidase 2 (gastrointestinal) (GPX2), mRNA
NM 002082	Homo sapiens G protein-coupled receptor kinase 6 (GPRK6), mRNA
NM 001504	Homo sapiens G protein-coupled receptor 9 (GPR9), mRNA
NM 001508	Homo sapiens G protein-coupled receptor 39 (GPR39), mRNA
NM 001507	Homo sapiens G protein-coupled receptor 38 (GPR38), mRNA
NM 001506	Homo sapiens G protein-coupled receptor 32 (GPR32), mRNA
NM 001505	Homo sapiens G protein-coupled receptor 30 (GPR30), mRNA
NM_001503	Homo sapiens glycosylphosphatidylinositol specific phospholipase D1 (GPLD1),
_	mRNA
NM_000408	Homo sapiens glycerol-3-phosphate dehydrogenase 2 (mitochondrial) (GPD2), mRNA
NM_001448	Homo sapiens glypican 4 (GPC4), mRNA
NM 002081	Homo sapiens glypican 1 (GPC1), mRNA
NM 000174	Homo sapiens glycoprotein IX (platelet) (GP9), mRNA
NM 000173	Homo sapiens glycoprotein Ib (platelet), alpha polypeptide (GP1BA), mRNA
NM_002080	Homo sapiens glutamic-oxaloacetic transaminase 2, mitochondrial (aspartate aminotransferase 2) (GOT2), nuclear gene encoding mitochondrial protein, mRNA
NM_002079	Homo sapiens glutamic-oxaloacetic transaminase 1, soluble (aspartate aminotransferase 1) (GOT1), mRNA
NM_002076	Homo sapiens glucosamine (N-acetyl)-6-sulfatase (Sanfilippo disease IIID) (GNS), mRNA
NM 001501	Homo sapiens gonadotropin-releasing hormone 2 (GNRH2), mRNA
NM_000825	Homo sapiens gonadotropin-releasing hormone 1 (leutinizing-releasing
	hormone) (GNRH1), mRNA
NM_002075	Homo sapiens guanine nucleotide binding protein (G protein), beta polypeptide 3
	(GNB3), mRNA
NM_002073	Homo sapiens guanine nucleotide binding protein (G protein), alpha z
ND 6 000170	polypeptide (GNAZ), mRNA Homo sapiens guanine nucleotide binding protein (G protein), alpha transducing
NM_000172	
NR 6 000070	activity polypeptide 1 (GNAT1), mRNA
NM_002072	Homo sapiens guanine nucleotide binding protein (G protein), q polypeptide (GNAQ), mRNA
NM_002071	Homo sapiens guanine nucleotide binding protein (G protein), alpha activating activity polypeptide, olfactory type (GNAL), mRNA
NM_002070	Homo sapiens guanine nucleotide binding protein (G protein), alpha inhibiting
	activity polypeptide 2 (GNAI2), mRNA
NM_002068	Homo sapiens guanine nucleotide binding protein (G protein), alpha 15 (Gq class) (GNA15), mRNA

201 00000	Homo sapiens guanine nucleotide binding protein (G protein), alpha 11 (Gq
NM_002067	class) (GNA11), mRNA
NM 003875	Homo sapiens guanine monphosphate synthetase (GMPS), mRNA
NM 002066	Homo sapiens GPI anchored molecule like protein (GML), mRNA
NM 001500	Homo sapiens GDP-mannose 4,6-dehydratase (GMDS), mRNA
NM 002065	Homo sapiens glutamate-ammonia ligase (glutamine synthase) (GLUL), mRNA
NM 002064	Homo sapiens glutaredoxin (thioltransferase) (GLRX), mRNA
NM 000824	Homo sapiens glycine receptor, beta (GLRB), mRNA
	Homo sapiens glycine receptor, alpha 2 (GLRA2), mRNA
NM_002063	Homo sapiens glucagon-like peptide 1 receptor (GLP1R), mRNA
NM_002062 NM_000170	Homo sapiens glycine dehydrogenase (decarboxylating; glycine decarboxylase,
14MT_000110	glycine cleavage system protein P) (GLDC), mRNA
NM 000169	Homo sapiens galactosidase, alpha (GLA), mRNA
NM 000167	Homo sapiens glycerol kinase (GK), mRNA
NM 000166	Homo sapiens gap junction protein, beta 1, 32kD (connexin 32, Charcot-Marie-
141AT_000100	Tooth neuropathy, X-linked) (GJB1), mRNA
NM 002060	Homo sapiens gap junction protein, alpha 4, 37kD (connexin 37) (GJA4), mRNA
NM 002060	Homo sapiens gastric inhibitory polypeptide receptor (GIPR), mRNA
NM 000164	Homo sapiens growth hormone releasing hormone receptor (GHRHR), mRNA
	Homo sapiens growth hormone receptor (GHR), mRNA
NM_000163 NM_000821	Homo sapiens gamma-glutamyl carboxylase (GGCX), mRNA
	Homo sapiens GDNF family receptor alpha 2 (GFRA2), mRNA
NM_001495	Homo sapiens GDNF family receptor alpha 2 (GFAP), intervi-
NM_002055	Homo sapiens genethonin 1 (GENX-3414), mRNA
NM 003943	Homo sapiens glial cell derived neurotrophic factor (GDNF), mRNA
NM_000514	Homo sapiens GDP dissociation inhibitor 1 (GDI1), mRNA
NM 001493	Homo sapiens GDF dissociation inhibitor 1 (GDF), inited at Homo sapiens glucosaminyl (N-acetyl) transferase 2, I-branching enzyme
NM_001491	(GCNT2), mRNA
NM 001490	Homo sapiens glucosaminyl (N-acetyl) transferase 1, core 2 (beta-1,6-N-
	acetylglucosaminyltransferase) (GCNT1), mRNA
NM 000160	Homo sapiens glucagon receptor (GCGR), mRNA
NM 002054	Homo sapiens glucagon (GCG), mRNA
NM 001485	Homo sapiens gastrulation brain homeo box 2 (GBX2), mRNA
NM 001483	Homo sapiens glioblastoma amplified sequence (GBAS), mRNA
NM 002048	Homo sapiens growth arrest-specific 1 (GAS1), mRNA
NM 001481	Homo sapiens growth arrest-specific 11 (GAS11), mRNA
NM 000819	Homo sapiens phosphoribosylglycinamide formyltransferase,
_	phosphoribosylglycinamide synthetase, phosphoribosylaminoimidazole
	synthetase (GART), mRNA
NM 002045	Homo sapiens growth associated protein 43 (GAP43), mRNA
NM 003614	Homo sapiens galanin receptor 3 (GALR3), mRNA
NM 000154	Homo sapiens galactokinase 1 (GALK1), mRNA
NM 001477	Homo sapiens G antigen 7B (GAGE7B), mRNA
NM_001476	Homo sapiens G antigen 6 (GAGE6), mRNA
NM 001475	Homo sapiens G antigen 5 (GAGE5), mRNA
NM 001474	Homo sapiens G antigen 4 (GAGE4), mRNA
NM 001473	Homo sapiens G antigen 3 (GAGE3), mRNA
NM 001472	Homo sapiens G antigen 2 (GAGE2), mRNA
NM 001468	Homo sapiens G antigen 1 (GAGE1), mRNA
NM_000818	Homo sapiens glutamate decarboxylase 2 (pancreatic islets and brain, 65kD)
NIM 002042	(GAD2), mRNA Homo sapiens gamma-aminobutyric acid (GABA) receptor, rho 2 (GABRR2),
NM_002043	1 Homo sapiens gamma-ammountyme acid (OADA) receptor, mo 2 (OADKR2),

	mRNA
NM 002042	Homo sapiens gamma-aminobutyric acid (GABA) receptor, rho 1 (GABRR1),
14141_002042	mRNA
NR (000402	Homo sapiens glucose-6-phosphate dehydrogenase (G6PD), nuclear gene
NM_000402	encoding mitochondrial protein, mRNA
ND (001460	Homo sapiens thyroid autoantigen 70kD (Ku antigen) (G22P1), mRNA
NM_001469	Homo sapiens FYN oncogene related to SRC, FGR, YES (FYN), mRNA
NM_002037	Homo sapiens Duffy blood group (FY), mRNA
NM_002036	Homo sapiens Dully blood group (F 1), likeva
NM_002035	Homo sapiens follicular lymphoma variant translocation 1 (FVT1), mRNA
NM_000150	Homo sapiens fucosyltransferase 6 (alpha (1,3) fucosyltransferase) (FUT6), mRNA
NM_002034	Homo sapiens fucosyltransferase 5 (alpha (1,3) fucosyltransferase) (FUT5), mRNA
NM_002033	Homo sapiens fucosyltransferase 4 (alpha (1,3) fucosyltransferase, myeloid-specific) (FUT4), mRNA
NM_000149	Homo sapiens fucosyltransferase 3 (galactoside 3(4)-L-fucosyltransferase, Lewis blood group included) (FUT3), mRNA
NM 000511	Homo sapiens fucosyltransferase 2 (secretor status included) (FUT2), mRNA
NM 000148	Homo sapiens fucosyltransferase 1 (galactoside 2-alpha-L-fucosyltransferase,
11112_000170	Bombay phenotype included) (FUT1), mRNA
NM 000147	Homo sapiens fucosidase, alpha-L- 1, tissue (FUCA1), mRNA
NM 002032	Homo sapiens ferritin, heavy polypeptide 1 (FTH1), mRNA
NM 000145	Homo sapiens follicle stimulating hormone receptor (FSHR), mRNA
NM 000510	Homo sapiens follicle stimulating hormone, beta polypeptide (FSHB), mRNA
NM 001463	Homo sapiens frizzled-related protein (FRZB), mRNA
NM 000144	Homo sapiens Friedreich ataxia (FRDA), mRNA
NM 001462	Homo sapiens formyl peptide receptor-like 1 (FPRL1), mRNA
NM 002029	Homo sapiens formyl peptide receptor 1 (FPR1), mRNA
NM 003838	Homo sapiens fucose-1-phosphate guanylyltransferase (FPGT), mRNA
NM 002027	Homo sapiens farnesyltransferase, CAAX box, alpha (FNTA), mRNA
	Homo sapiens fragile X mental retardation 2 (FMR2), mRNA
NM 002025	Homo sapiens fragile X mental retardation 1 (FMR1), mRNA
NM 002024	Homo sapiens flavin containing monooxygenase 5 (FMO5), mRNA
NM_001461	Homo sapiens flavin containing monooxygenase 4 (FMO4), mRNA
NM_002022	Homo sapiens flavin containing monooxygenase 2 (FMO2), mRNA
NM_001460	Homo sapiens flavin containing monooxygenase 2 (FMO2), mRNA
NM_002021	Homo sapiens flavili containing monooxygenase i (FWO), meters
NM_002020	Homo sapiens fms-related tyrosine kinase 4 (FLT4), mRNA Homo sapiens fms-related tyrosine kinase 3 ligand (FLT3LG), mRNA
NM_001459	Homo sapiens ims-related tyrosine kinase 5 ligand (PLISCO), index
NM_002019	Homo sapiens fms-related tyrosine kinase 1 (vascular endothelial growth
27.5.632.155	factor/vascular permeability factor receptor) (FLT1), mRNA
NM_001455_	Homo sapiens forkhead box O3A (FOXO3A), mRNA
NM_001453	Homo sapiens forkhead box C1 (FOXC1), mRNA
NM_001451	Homo sapiens forkhead box F1 (FOXF1), mRNA
NM_001450	Homo sapiens four and a half LIM domains 2 (FHL2), mRNA
NM_001449	Homo sapiens four and a half LIM domains 1 (FHL1), mRNA
NM_002012	Homo sapiens fragile histidine triad gene (FHIT), mRNA
NM_000143	Homo sapiens fumarate hydratase (FH), mRNA
NM_002002	Homo sapiens Fc fragment of IgE, low affinity II, receptor for (CD23A) (FCER2), mRNA
NM_002001	Homo sapiens Fc fragment of IgE, high affinity I, receptor for; alpha polypeptide (FCER1A), mRNA
NM 002000	Homo sapiens Fc fragment of IgA, receptor for (FCAR), mRNA

NM_003837	Homo sapiens fructose-1,6-bisphosphatase 2 (FBP2), mRNA
NM_001998	Homo sapiens fibulin 2 (FBLN2), mRNA
NM 003923	Homo sapiens forkhead box H1 (FOXH1), mRNA
NM 003950	Homo sapiens coagulation factor II (thrombin) receptor-like 3 (F2RL3), mRNA
NM 003975	Homo sapiens SH2 domain protein 2A (SH2D2A), mRNA
NM 001440	Homo sapiens exostoses (multiple)-like 3 (EXTL3), mRNA
NM 001988	Homo sapiens envoplakin (EVPL), mRNA
NM 001985	Homo sapiens electron-transfer-flavoprotein, beta polypeptide (ETFB), mRNA
NM_000126	Homo sapiens electron-transfer-flavoprotein, alpha polypeptide (glutaric aciduria II) (ETFA), nuclear gene encoding mitochondrial protein, mRNA
NM 001438	Homo sapiens estrogen-related receptor gamma (ESRRG), mRNA
NM 000125	Homo sapiens estrogen receptor 1 (ESR1), mRNA
NM 000123	Homo sapiens excision repair cross-complementing rodent repair deficiency,
14141_000123	complementation group 5 (xeroderma pigmentosum, complementation group G
	(Cockayne syndrome)) (ERCC5), mRNA
NM_001983	Homo sapiens excision repair cross-complementing rodent repair deficiency,
1414_001505	complementation group 1 (includes overlapping antisense sequence) (ERCC1), mRNA
NB4 000502	Homo sapiens eosinophil peroxidase (EPX), mRNA
NM_000502	Homo sapiens epidermal growth factor receptor pathway substrate 15 (EPS15),
NM_001981	mRNA
NM_000799	Homo sapiens erythropoietin (EPO), mRNA
NM_001980	Homo sapiens epimorphin (EPIM), mRNA
NM_001431	Homo sapiens erythrocyte membrane protein band 4.1-like 2 (EPB41L2), mRNA
NM_001430	Homo sapiens endothelial PAS domain protein 1 (EPAS1), mRNA
NM_001977	Homo sapiens glutamyl aminopeptidase (aminopeptidase A) (ENPEP), mRNA
NM_001974	Homo sapiens egf-like module containing, mucin-like, hormone receptor-like sequence 1 (EMR1), mRNA
NM 001425	Homo sapiens epithelial membrane protein 3 (EMP3), mRNA
NM 001424	Homo sapiens epithelial membrane protein 2 (EMP2), mRNA
NM 001423	Homo sapiens epithelial membrane protein 1 (EMP1), mRNA
NM 001421	Homo sapiens E74-like factor 4 (ets domain transcription factor) (ELF4), mRNA
NM_001419	Homo sapiens ELAV (embryonic lethal, abnormal vision, Drosophila)-like 1 (Hu antigen R) (ELAVL1), mRNA
NM 001972	Homo sapiens elastase 2, neutrophil (ELA2), mRNA
NM 001972	Homo sapiens eukaryotic translation initiation factor 5A (EIF5A), mRNA
	Homo sapiens eukaryotic translation initiation factor 4 gamma, 2 (EIF4G2),
NM_001418	mRNA
NM_003732	Homo sapiens eukaryotic translation initiation factor 4E binding protein 3 (EIF4EBP3), mRNA
NM_001968	Homo sapiens eukaryotic translation initiation factor 4E (EIF4E), mRNA
NM_001416	Homo sapiens eukaryotic translation initiation factor 4A, isoform 1 (EIF4A1), mRNA
NM_003753	Homo sapiens eukaryotic translation initiation factor 3, subunit 7 (zeta, 66/67kD) (EIF3S7), mRNA
NM_001568	Homo sapiens eukaryotic translation initiation factor 3, subunit 6 (48kD) (EIF3S6), mRNA
NM_003754	Homo sapiens eukaryotic translation initiation factor 3, subunit 5 (epsilon, 47kD)
11.1_000,04	(EIF3S5), mRNA
NM_003757	Homo sapiens eukaryotic translation initiation factor 3, subunit 2 (beta, 36kD) (EIF3S2), mRNA
NM_003750	Homo sapiens eukaryotic translation initiation factor 3, subunit 10 (theta,

	150/170kD) (EIF3S10), mRNA
NM_001415	Homo sapiens eukaryotic translation initiation factor 2, subunit 3 (gamma, 52kD)
14101_001413	(EIF2S3), mRNA
NM_003908	Homo sapiens eukaryotic translation initiation factor 2, subunit 2 (beta, 38kD)
14141_002308	(EIF2S2), mRNA
NM 001966	Homo sapiens enoyl-Coenzyme A, hydratase/3-hydroxyacyl Coenzyme A
14147_001300	dehydrogenase (EHHADH), nuclear gene encoding mitochondrial protein,
	mRNA
NM 001965	Homo sapiens early growth response 4 (EGR4), mRNA
NM 001964	Homo sapiens early growth response 1 (EGR1), mRNA
NM 001406	Homo sapiens ephrin-B3 (EFNB3), mRNA
NM 001962	Homo sapiens ephrin-A5 (EFNA5), mRNA
NM 001405	Homo sapiens ephrin-A2 (EFNA2), mRNA
NM 001961	Homo sapiens eukaryotic translation elongation factor 2 (EEF2), mRNA
NM 001958	Homo sapiens eukaryotic translation elongation factor 1 alpha 2 (EEF1A2),
	mRNA
NM_001956	Homo sapiens endothelin 2 (EDN2), mRNA
NM_001955	Homo sapiens endothelin 1 (EDN1), mRNA
NM_003775	Homo sapiens endothelial differentiation, G-protein-coupled receptor 6 (EDG6),
	mRNA
NM_001399	Homo sapiens ectodermal dysplasia 1, anhidrotic (ED1), mRNA
NM_001397	Homo sapiens endothelin converting enzyme 1 (ECE1), mRNA
NM_003240	Homo sapiens endometrial bleeding associated factor (left-right determination,
	factor A; transforming growth factor beta superfamily) (EBAF), mRNA
NM_001948	Homo sapiens dUTP pyrophosphatase (DUT), mRNA
NM_001945	Homo sapiens diphtheria toxin receptor (heparin-binding epidermal growth
	factor-like growth factor) (DTR), mRNA
NM_001939	Homo sapiens dystrophin related protein 2 (DRP2), mRNA
NM_001938	Homo sapiens down-regulator of transcription 1, TBP-binding (negative cofactor 2) (DR1), mRNA
NM 001387	Homo sapiens dihydropyrimidinase-like 3 (DPYSL3), mRNA
NM 001385	Homo sapiens dihydropyrimidinase (DPYS), mRNA
NM 001935	Homo sapiens dipeptidylpeptidase IV (CD26, adenosine deaminase complexing
	protein 2) (DPP4), mRNA
NM_003863	Homo sapiens dolichyl-phosphate mannosyltransferase polypeptide 2, regulatory subunit (DPM2), mRNA
NM 001380	Homo sapiens dedicator of cyto-kinesis 1 (DOCK1), mRNA
NM_001379	Homo sapiens DNA (cytosine-5-)-methyltransferase 1 (DNMT1), mRNA
NM 001375	Homo sapiens deoxyribonuclease II, lysosomal (DNASE2), mRNA
NM 001374	Homo sapiens deoxyribonuclease I, lysosomar (Britisba), mRNA
NM 001934	Homo sapiens distal-less homeobox 4 (DLX4), mRNA
NM 001933	Homo sapiens dihydrolipoamide S-succinyltransferase (E2 component of 2-oxo-
1444_001733	glutarate complex) (DLST), mRNA
NM 001362	Homo sapiens deiodinase, iodothyronine, type III (DIO3), mRNA
NM 001360	Homo sapiens 7-dehydrocholesterol reductase (DHCR7), mRNA
NM_003670	Homo sapiens basic helix-loop-helix domain containing, class B, 2 (BHLHB2),
NR 001271	mRNA However and a last and action for its 1 member C2 (dibydrodio)
NM_001354	Homo sapiens aldo-keto reductase family 1, member C2 (dihydrodiol
	dehydrogenase 2; bile acid binding protein; 3-alpha hydroxysteroid
ND (000700	dehydrogenase, type III) (AKR1C2), mRNA Homo sapiens dopa decarboxylase (aromatic L-amino acid decarboxylase)
NM_000790	(DDC), mRNA
L	(DDO), HILLIAN

NM_000789	Homo sapiens dipeptidyl carboxypeptidase 1 (angiotensin I converting enzyme)
27.6 204000	(ACE), mRNA
NM_001920	Homo sapiens decorin (DCN), mRNA
NM_000788	Homo sapiens deoxycytidine kinase (DCK), mRNA
NM_001919	Homo sapiens dodecenoyl-Coenzyme A delta isomerase (3,2 trans-enoyl-Coenzyme A isomerase) (DCI), mRNA
NM_001918	Homo sapiens dihydrolipoamide branched chain transacylase (E2 component of branched chain keto acid dehydrogenase complex; maple syrup urine disease) (DBT), mRNA
NM_001352	Homo sapiens D site of albumin promoter (albumin D-box) binding protein (DBP), mRNA
NM 001351	Homo sapiens deleted in azoospermia-like (DAZL), mRNA
NM 001350	Homo sapiens death-associated protein 6 (DAXX), mRNA
NM 001344	Homo sapiens defender against cell death 1 (DAD1), mRNA
NM 003472	Homo sapiens DEK oncogene (DNA binding) (DEK), mRNA
NM_000776	Homo sapiens cytochrome P450, subfamily IIIA (niphedipine oxidase), polypeptide 3 (CYP3A3), mRNA
NM 001916	Homo sapiens cytochrome c-1 (CYC1), mRNA
NM_001914	Homo sapiens cytochrome b-5 (CYB5), nuclear gene encoding mitochondrial protein, mRNA
NM 003928	Homo sapiens CAAX box 1 (CXX1), mRNA
NM 003611	Homo sapiens chromosome X open reading frame 5 (CXORF5), mRNA
NM 003467	Homo sapiens chemokine (C-X-C motif), receptor 4 (fusin) (CXCR4), mRNA
NM 001338	Homo sapiens coxsackie virus and adenovirus receptor (CXADR), mRNA
NM 003478	Homo sapiens cullin 5 (CUL5), mRNA
NM 003591	Homo sapiens cullin 2 (CUL2), mRNA
NM 001336	Homo sapiens cathepsin Z (CTSZ), mRNA
NM 001335	Homo sapiens cathepsin W (lymphopain) (CTSW), mRNA
NM 001912	Homo sapiens cathepsin L (CTSL), mRNA
	Homo sapiens cathepsin L2 (CTSL2), mRNA
NM_001333	Homo sapiens cathepsin K (pycnodysostosis) (CTSK), mRNA
NM_000396	Homo sapiens cathepsin K (pychodysostosis) (CTSK), micryk Homo sapiens cathepsin G (CTSG), mRNA
NM_001911	
NM_001910	Homo sapiens cathepsin E (CTSE), mRNA
NM_001909	Homo sapiens cathepsin D (lysosomal aspartyl protease) (CTSD), mRNA
NM_001814	Homo sapiens cathepsin C (CTSC), mRNA
NM_001908	Homo sapiens cathepsin B (CTSB), mRNA
NM_001907	Homo sapiens chymotrypsin-like (CTRL), mRNA
NM_001906	Homo sapiens chymotrypsinogen B1 (CTRB1), mRNA
NM_001905	Homo sapiens CTP synthase (CTPS), mRNA
NM_001904	Homo sapiens catenin (cadherin-associated protein), beta 1 (88kD) (CTNNB1), mRNA
NM_003798	Homo sapiens catenin (cadherin-associated protein), alpha-like 1 (CTNNAL1), mRNA
NM_001903	Homo sapiens catenin (cadherin-associated protein), alpha 1 (102kD) (CTNNA1), mRNA
NM 001902	Homo sapiens cystathionase (cystathionine gamma-lyase) (CTH), mRNA
NM 001901	Homo sapiens connective tissue growth factor (CTGF), mRNA
NM 001330	Homo sapiens cardiotrophin 1 (CTF1), mRNA
NM 000100	Homo sapiens cystatin B (stefin B) (CSTB), mRNA
NM 003650	Homo sapiens cystatin F (leukocystatin) (CST7), mRNA
NM_001323	Homo sapiens cystatin F/M (CST6), mRNA
NM 001900	Homo sapiens cystatin D (CST5), mRNA
T4141 001300	Tronic septons cysican D (CS12), micris

NR 4 001000	Homo sapiens cystatin S (CST4), mRNA
NM_001899	Homo sapiens cystatin S (CST4), inktvA Homo sapiens cystatin C (amyloid angiopathy and cerebral hemorrhage) (CST3),
NM_000099	
	mRNA
NM_001322	Homo sapiens cystatin SA (CST2), mRNA
NM_001898	Homo sapiens cystatin SN (CST1), mRNA
NM_001321	Homo sapiens cysteine and glycine-rich protein 2 (CSRP2), mRNA
NM_001896	Homo sapiens casein kinase 2, alpha prime polypeptide (CSNK2A2), mRNA
NM_001895	Homo sapiens casein kinase 2, alpha 1 polypeptide (CSNK2A1), mRNA
NM_001894	Homo sapiens casein kinase 1, epsilon (CSNK1E), mRNA
NM_001893	Homo sapiens casein kinase 1, delta (CSNK1D), mRNA
NM_001892	Homo sapiens casein kinase 1, alpha 1 (CSNK1A1), mRNA
NM_001891	Homo sapiens casein, beta (CSN2), mRNA
NM_001890	Homo sapiens casein, alpha (CSN1), mRNA
NM_000760	Homo sapiens colony stimulating factor 3 receptor (granulocyte) (CSF3R),
	mRNA
NM_000759	Homo sapiens colony stimulating factor 3 (granulocyte) (CSF3), mRNA
NM_000758	Homo sapiens colony stimulating factor 2 (granulocyte-macrophage) (CSF2),
-	mRNA
NM_000757	Homo sapiens colony stimulating factor 1 (macrophage) (CSF1), mRNA
NM_003651	Homo sapiens cold shock domain protein A (CSDA), mRNA
NM_001315	Homo sapiens mitogen-activated protein kinase 14 (MAPK14), mRNA
NM_001884	Homo sapiens cartilage linking protein 1 (CRTL1), mRNA
NM_001313	Homo sapiens collapsin response mediator protein 1 (CRMP1), mRNA
NM 001312	Homo sapiens cysteine-rich protein 2 (CRIP2), mRNA
NM 001311	Homo sapiens cysteine-rich protein 1 (intestinal) (CRIP1), mRNA
NM 000756	Homo sapiens corticotropin releasing hormone (CRH), mRNA
NM 001881	Homo sapiens cAMP responsive element modulator (CREM), mRNA
NM 003851	Homo sapiens cellular repressor of E1A-stimulated genes (CREG), mRNA
NM_001310	Homo sapiens cAMP responsive element binding protein-like 2 (CREBL2), mRNA
NM 001880	Homo sapiens activating transcription factor 2 (ATF2), mRNA
NM_003805	Homo sapiens CASP2 and RIPK1 domain containing adaptor with death domain (CRADD), mRNA
NM_001877	Homo sapiens complement component (3d/Epstein Barr virus) receptor 2 (CR2),
NM 000098	mRNA Homo sapiens carnitine palmitoyltransferase II (CPT2), nuclear gene encoding
74747_000038	mitochondrial protein, mRNA
NM_001876	Homo sapiens carnitine palmitoyltransferase I, liver (CPT1A), nuclear gene
TATAT OO 1910	encoding mitochondrial protein, mRNA
NM 001875	Homo sapiens carbamoyl-phosphate synthetase 1, mitochondrial (CPS1), nuclear
14147_001012	gene encoding mitochondrial protein, mRNA
NM 000097	Homo sapiens coproporphyrinogen oxidase (coproporphyria, harderoporphyria)
14747_000027	(CPO), mRNA
NM 001871	Homo sapiens carboxypeptidase B1 (tissue) (CPB1), mRNA
NM 001870	Homo sapiens carboxypeptidase A3 (mast cell) (CPA3), mRNA
NM 001869	Homo sapiens carboxypeptidase A2 (pancreatic) (CPA2), mRNA
NM 001868	Homo sapiens carboxypeptidase A2 (pancreatic) (CPA1), mRNA
NM 001808	Homo sapiens beaded filament structural protein 2, phakinin (BFSP2), mRNA
NM 001302	Homo sapiens cortistatin (CORT), mRNA
NM 001302 NM 003832	Homo sapiens cortistatif (CORT), fileNA Homo sapiens phosphoserine phosphatase-like (PSPHL), mRNA
NM 001843	Homo sapiens contactin 1 (CNTN1), mRNA
NM 001843	Homo sapiens contactin 1 (CNTN1), interval Homo sapiens ciliary neurotrophic factor rec ptor (CNTFR), mRNA
19191 UU 1 042	1 Monte suprems emany neuroproprie factor for port (Chilles), marri-

37 C 001020	Homo sapiens calponin 3, acidic (CNN3), mRNA
NM_001839	Homo sapiens calponin 1, basic, smooth muscle (CNN1), mRNA
NM 001299	Homo sapiens cyclic nucleotide gated channel beta 1 (CNGB1), mRNA
NM_001297	Homo sapiens cyclic nucleotide gated channel alpha 3 (CNGA3), mRNA
NM 001298	Homo sapiens cyclic nucleotide gated channel alpha 1 (CNGA1), mRNA
NM_000087	Homo sapiens cyclic nucleotide gated channel alpha I (CNOAI), mid A Homo sapiens cytidine monophosphate-N-acetylneuraminic acid hydroxylase
NM_003570	(CMP-N-acetylneuraminate monooxygenase) (CMAH), mRNA
27.6.001026	Homo sapiens chymase 1, mast cell (CMA1), mRNA
NM_001836	Homo sapiens chymase 1, mast cen (CMA1), matter Homo sapiens clusterin (complement lysis inhibitor, SP-40,40, sulfated
NM_001831	glycoprotein 2, testosterone-repressed prostate message 2, apolipoprotein J)
	(CLU), mRNA
NM_001294	Homo sapiens cleft lip and palate associated transmembrane protein 1
14141_001294	(CLPTM1), mRNA
NM_003476	Homo sapiens cysteine and glycine-rich protein 3 (cardiac LIM protein)
14141_005470	(CSRP3), mRNA
NM_001293	Homo sapiens chloride channel, nucleotide-sensitive, 1A (CLNS1A), mRNA
NM_003277	Homo sapiens claudin 5 (transmembrane protein deleted in velocardiofacial
	syndrome) (CLDN5), mRNA
NM 001306	Homo sapiens claudin 3 (CLDN3), mRNA
NM 001829	Homo sapiens chloride channel 3 (CLCN3), mRNA
NM_001284	Homo sapiens adaptor-related protein complex 3, sigma 1 subunit (AP3S1),
_	mRNA
NM 001827	Homo sapiens CDC28 protein kinase 2 (CKS2), mRNA
NM 001826	Homo sapiens CDC28 protein kinase 1 (CKS1), mRNA
NM 001824	Homo sapiens creatine kinase, muscle (CKM), mRNA
NM 001823	Homo sapiens creatine kinase, brain (CKB), mRNA
NM 001281	Homo sapiens cytoskeleton-associated protein 1 (CKAP1), mRNA
NM 003613	Homo sapiens cartilage intermediate layer protein, nucleotide
	pyrophosphohydrolase (CILP), mRNA
NM_001278	Homo sapiens conserved helix-loop-helix ubiquitous kinase (CHUK), mRNA
NM_003654	Homo sapiens carbohydrate (chondroitin 6/keratan) sulfotransferase 1 (CHST1),
	mRNA
NM_000750	Homo sapiens cholinergic receptor, nicotinic, beta polypeptide 4 (CHRNB4),
ND (000740	mRNA Homo sapiens cholinergic receptor, nicotinic, beta polypeptide 3 (CHRNB3),
NM_000749	mRNA
NM 000748	Homo sapiens cholinergic receptor, nicotinic, beta polypeptide 2 (neuronal)
14147_000/40	(CHRNB2), mRNA
NM 000746	Homo sapiens cholinergic receptor, nicotinic, alpha polypeptide 7 (CHRNA7),
1111_000740	mRNA
NM 000745	Homo sapiens cholinergic receptor, nicotinic, alpha polypeptide 5 (CHRNA5),
	mRNA
NM 000744	Homo sapiens cholinergic receptor, nicotinic, alpha polypeptide 4 (CHRNA4),
	mRNA
NM 000743	Homo sapiens cholinergic receptor, nicotinic, alpha polypeptide 3 (CHRNA3),
	mRNA
NM 000742	Homo sapiens cholinergic receptor, nicotinic, alpha polypeptide 2 (neuronal)
	(CHRNA2), mRNA
NM_000741	Homo sapiens cholinergic receptor, muscarinic 4 (CHRM4), mRNA
NM_000740	Homo sapiens cholinergic receptor, muscarinic 3 (CHRM3), mRNA
NM_000739	Homo sapiens cholinergic receptor, muscarinic 2 (CHRM2), mRNA
NM_000738	Homo sapiens cholinergic receptor, muscarinic 1 (CHRM1), mRNA

NM 001822	Homo sapiens chimerin (chimaerin) 1 (CHN1), mRNA
NM 001821	Homo sapiens choroideremia-like (Rab escort protein 2) (CHML), mRNA
NM 001819	Homo sapiens chromogranin B (secretogranin 1) (CHGB), mRNA
NM 001269	Homo sapiens chromosome condensation 1 (CHC1), mRNA
NM 001267	Homo sapiens chondroadherin (CHAD), mRNA
NM_001817	Homo sapiens carcinoembryonic antigen-related cell adhesion molecule 4
_	(CEACAM4), mRNA
NM_001816	Homo sapiens carcinoembryonic antigen-related cell adhesion molecule 8
-	(CEACAM8), mRNA
NM_001815	Homo sapiens carcinoembryonic antigen-related cell adhesion molecule 3
-	(CEACAM3), mRNA
NM_003663	Homo sapiens CGG triplet repeat binding protein 1 (CGGBP1), mRNA
NM_001813	Homo sapiens centromere protein E (312kD) (CENPE), mRNA
NM_001808	Homo sapiens carboxyl ester lipase-like (bile salt-stimulated lipase-like) (CELL),
-	mRNA
NM_001807	Homo sapiens carboxyl ester lipase (bile salt-stimulated lipase) (CEL), mRNA
NM_001805	Homo sapiens CCAAT/enhancer binding protein (C/EBP), epsilon (CEBPE),
	mRNA CODYO DATA
NM_001265	Homo sapiens caudal type homeo box transcription factor 2 (CDX2), mRNA
NM_001804	Homo sapiens caudal type homeo box transcription factor 1 (CDX1), mRNA
NM_001803	Homo sapiens CDW52 antigen (CAMPATH-1 antigen) (CDW52), mRNA
NM_001264	Homo sapiens corneodesmosin (CDSN), mRNA
NM_001263	Homo sapiens CDP-diacylglycerol synthase (phosphatidate cytidylyltransferase)
	1 (CDS1), mRNA
NM_001801	Homo sapiens cysteine dioxygenase, type I (CDO1), mRNA
NM_001769	Homo sapiens CD9 antigen (p24) (CD9), mRNA
NM_001768	Homo sapiens CD8 antigen, alpha polypeptide (p32) (CD8A), mRNA
NM_003874	Homo sapiens CD84 antigen (leukocyte antigen) (CD84), mRNA
NM_001781	Homo sapiens CD69 antigen (p60, early T-cell activation antigen) (CD69),
ND (001790	mRNA Homo sapiens CD63 antigen (melanoma 1 antigen) (CD63), mRNA
NM_001780	Homo sapiens CD58 antigen (Inclandina 1 antigen) (GD56), including Homo sapiens CD58 antigen, (lymphocyte function-associated antigen 3)
NM_001779	(CD58), mRNA
NM 001778	Homo sapiens CD48 antigen (B-cell membrane protein) (CD48), mRNA
NM_001777	Homo sapiens CD47 antigen (Rh-related antigen, integrin-associated signal
	transducer) (CD47), mRNA
NM_000733	Homo sapiens CD3E antigen, epsilon polypeptide (TiT3 complex) (CD3E), mRNA
NM_000732	Homo sapiens CD3D antigen, delta polypeptide (TiT3 complex) (CD3D),
1	mRNA
NM 001776	Homo sapiens ectonucleoside triphosphate diphosphohydrolase 1 (ENTPD1),
	mRNA
NM_001775	Homo sapiens CD38 antigen (p45) (CD38), mRNA
NM_001774	Homo sapiens CD37 antigen (CD37), mRNA
NM 001773	Homo sapiens CD34 antigen (CD34), mRNA
NM 003830	Homo sapiens sialic acid binding Ig-like lectin 5 (SIGLEC5), mRNA
NM 001245	Homo sapiens sialic acid binding Ig-like lectin 6 (SIGLEC6), mRNA
NM 001772	Homo sapiens CD33 antigen (gp67) (CD33), mRNA
NM 001767	Homo sapiens CD2 antigen (p50), sheep red blood cell receptor (CD2), mRNA
NM 001771	Homo sapiens CD22 antigen (CD22), mRNA
NM 001766	Homo sapiens CD1D antigen, d polypeptide (CD1D), mRNA
NM 001765	Homo sapiens CD1C antigen, c polypeptide (CD1C), mRNA

	CDID wise hashwaride (CDID) mPNA
NM_001764	Homo sapiens CD1B antigen, b polypeptide (CD1B), mRNA
NM_001838	Homo sapiens chemokine (C-C motif) receptor 7 (CCR7), mRNA
NM_001837	Homo sapiens chemokine (C-C motif) receptor 3 (CCR3), mRNA
NM_001758	Homo sapiens cyclin D1 (PRAD1 parathyroid adenomatosis 1) (CCND1),
	mRNA
NM_000731	Homo sapiens cholecystokinin B receptor (CCKBR), mRNA
NM_000730	Homo sapiens cholecystokinin A receptor (CCKAR), mRNA
NM_001757	Homo sapiens carbonyl reductase 1 (CBR1), mRNA
NM_001754	Homo sapiens runt-related transcription factor 1 (acute myeloid leukemia 1; aml1 oncogene) (RUNX1), mRNA
NM_003688	Homo sapiens calcium/calmodulin-dependent serine protein kinase (MAGUK family) (CASK), mRNA
NM 001747	Homo sapiens capping protein (actin filament), gelsolin-like (CAPG), mRNA
NM_001744	Homo sapiens calcium/calmodulin-dependent protein kinase IV (CAMK4), mRNA
NM_001743	Homo sapiens calmodulin 2 (phosphorylase kinase, delta) (CALM2), mRNA
NM 001742	Homo sapiens calcitonin receptor (CALCR), mRNA
NM 001741	Homo sapiens calcitonin/calcitonin-related polypeptide, alpha (CALCA), mRNA
NM_000727	Homo sapiens calcium channel, voltage-dependent, gamma subunit l
	(CACNG1), mRNA
NM_000726	Homo sapiens calcium channel, voltage-dependent, beta 4 subunit (CACNB4), mRNA
NM_000725	Homo sapiens calcium channel, voltage-dependent, beta 3 subunit (CACNB3), mRNA
NM_000724	Homo sapiens calcium channel, voltage-dependent, beta 2 subunit (CACNB2), mRNA
NM_000723	Homo sapiens calcium channel, voltage-dependent, beta 1 subunit (CACNB1), mRNA
NM_000721	Homo sapiens calcium channel, voltage-dependent, alpha 1E subunit (CACNA1E), mRNA
NM_000720	Homo sapiens calcium channel, voltage-dependent, L type, alpha 1D subunit (CACNA1D), mRNA
NM_000719	Homo sapiens calcium channel, voltage-dependent, L type, alpha 1C subunit (CACNA1C), mRNA
NM_000718	Homo sapiens calcium channel, voltage-dependent, L type, alpha 1B subunit (CACNA1B), mRNA
NM_001739	Homo sapiens carbonic anhydrase VA, mitochondrial (CA5A), nuclear gene encoding mitochondrial protein, mRNA
NM 001738	Homo sapiens carbonic anhydrase I (CA1), mRNA
NM_001737	Homo sapiens complement component 9 (C9), mRNA
NM 001736	Homo sapiens complement component 5 receptor 1 (C5a ligand) (C5R1), mRNA
NM 001735	Homo sapiens complement component 5 (C5), mRNA
NM 003956	Homo sapiens cholesterol 25-hydroxylase (CH25H), mRNA
NM_001734	Homo sapiens complement component 1, s subcomponent (C1S), mRNA
NM_001733	Homo sapiens complement component 1, r subcomponent (C1R), mRNA
NM 001732	Homo sapiens butyrophilin, subfamily 1, member A1 (BTN1A1), mRNA
NM 001731	Homo sapiens B-cell translocation gene 1, anti-proliferative (BTG1), mRNA
NM 001729	Homo sapiens betacellulin (BTC), mRNA
NM 001728	Homo sapiens basigin (BSG), mRNA
NM_003742	Homo sapiens ATP-binding cassette, sub-family B (MDR/TAP), member 11 (ABCB11), mRNA
NM 001727	Homo sapiens bombesin-like receptor 3 (BRS3), mRNA

NM 000059	Homo sapiens breast cancer 2, early onset (BRCA2), mRNA
NM 001725	Homo sapiens bactericidal/permeability-increasing protein (BPI), mRNA
NM 001724	Homo sapiens 2,3-bisphosphoglycerate mutase (BPGM), mRNA
NM 001723	Homo sapiens bullous pemphigoid antigen 1 (230/240kD) (BPAG1), mRNA
NM 001717	Homo sapiens basonuclin (BNC), mRNA
NM 001722	Homo sapiens BN51 (BHK21) temperature sensitivity complementing (BN51T),
11112_00112=	mRNA
NM 001721	Homo sapiens BMX non-receptor tyrosine kinase (BMX), mRNA
NM 001203	Homo sapiens bone morphogenetic protein receptor, type IB (BMPR1B), mRNA
NM 001720	Homo sapiens bone morphogenetic protein 8 (osteogenic protein 2) (BMP8),
	mRNA
NM 001719	Homo sapiens bone morphogenetic protein 7 (osteogenic protein 1) (BMP7),
	mRNA
NM 001202	Homo sapiens bone morphogenetic protein 4 (BMP4), mRNA
NM 000713	Homo sapiens biliverdin reductase B (flavin reductase (NADPH)) (BLVRB),
_	mRNA
NM_000712	Homo sapiens biliverdin reductase A (BLVRA), mRNA
NM_001713	Homo sapiens betaine-homocysteine methyltransferase (BHMT), mRNA
NM_001712	Homo sapiens carcinoembryonic antigen-related cell adhesion molecule 1
	(biliary glycoprotein) (CEACAM1), mRNA
NM_001711	Homo sapiens biglycan (BGN), mRNA
NM_000711	Homo sapiens bone gamma-carboxyglutamate (gla) protein (osteocalcin)
	(BGLAP), mRNA
NM_001709	Homo sapiens brain-derived neurotrophic factor (BDNF), mRNA
NM_000710	Homo sapiens bradykinin receptor B1 (BDKRB1), mRNA
NM_001707	Homo sapiens B-cell CLL/lymphoma 7B (BCL7B), mRNA
NM_001706	Homo sapiens B-cell CLL/lymphoma 6 (zinc finger protein 51) (BCL6), mRNA
NM_003921	Homo sapiens B-cell CLL/lymphoma 10 (BCL10), mRNA
NM_003657	Homo sapiens breast carcinoma amplified sequence 1 (BCAS1), mRNA
NM_001188	Homo sapiens BCL2-antagonist/killer 1 (BAK1), mRNA
NM_001704	Homo sapiens brain-specific angiogenesis inhibitor 3 (BAI3), mRNA
NM_001703	Homo sapiens brain-specific angiogenesis inhibitor 2 (BAI2), mRNA
NM_001702	Homo sapiens brain-specific angiogenesis inhibitor 1 (BAI1), mRNA
NM_001186	Homo sapiens BTB and CNC homology 1, basic leucine zipper transcription
	factor 1 (BACH1), mRNA
NM_001701	Homo sapiens bile acid Coenzyme A amino acid N-acyltransferase (glycine N-
	choloyltransferase) (BAAT), mRNA
NM_001185	Homo sapiens alpha-2-glycoprotein 1, zinc (AZGP1), mRNA
NM_001184	Homo sapiens ataxia telangiectasia and Rad3 related (ATR), mRNA
NM_000053	Homo sapiens ATPase, Cu++ transporting, beta polypeptide (Wilson disease)
27.6 600015	(ATP7B), mRNA
NM_003945	Homo sapiens ATPase, H+ transporting, lysosomal (vacuolar proton pump) 9kD
ND 5 001 606	(ATP6H), mRNA
NM_001696	Homo sapiens ATPase, H+ transporting, lysosomal (vacuolar proton pump)
ND (001 (00	31kD (ATP6E), mRNA
NM_001693	Homo sapiens ATPase, H+ transporting, lysosomal (vacuolar proton pump), beta
ND 6 001 (02	polypeptide, 56/58kD, isoform 2 (ATP6B2), mRNA
NM_001692	Homo sapiens ATPase, H+ transporting, lysosomal (vacuolar proton pump), beta
NB/ 001/01	polypeptide, 56/58kD, isoform 1 (ATP6B1), mRNA
NM_001691	Homo sapiens ATPase, H+ transporting, lysosomal (vacuolar proton pump),
NR4 001600	alpha polypeptide, 70kD, isoform 2 (ATP6A2), mRNA
NM_001690	Homo sapiens ATPase, H+ transporting, lysosomal (vacuolar proton pump),

	11 1 ACC TOLD :- C 1 (ATDCA1) DNA
	alpha polypeptide, 70kD, isoform 1 (ATP6A1), mRNA
NM_001697	Homo sapiens ATP synthase, H+ transporting, mitochondrial F1 complex, O
	subunit (oligomycin sensitivity conferring protein) (ATP50), mRNA
NM_001686	Homo sapiens ATP synthase, H+ transporting, mitochondrial F1 complex, beta
	polypeptide (ATP5B), nuclear gene encoding mitochondrial protein, mRNA
NM_000704	Homo sapiens ATPase, H+/K+ exchanging, alpha polypeptide (ATP4A), mRNA
NM_001684	Homo sapiens ATPase, Ca++ transporting, plasma membrane 4 (ATP2B4), mRNA
NM_001682	Homo sapiens ATPase, Ca++ transporting, plasma membrane 1 (ATP2B1), mRNA
NM_001681	Homo sapiens ATPase, Ca++ transporting, cardiac muscle, slow twitch 2 (ATP2A2), mRNA
NM_001679	Homo sapiens ATPase, Na+/K+ transporting, beta 3 polypeptide (ATP1B3), mRNA
NM_001678	Homo sapiens ATPase, Na+/K+ transporting, beta 2 polypeptide (ATP1B2), mRNA
NM_001677	Homo sapiens ATPase, Na+/K+ transporting, beta 1 polypeptide (ATP1B1), mRNA
NM_000703	Homo sapiens ATPase, Na+/K+ transporting, alpha 3 polypeptide (ATP1A3), mRNA
NM_000702	Homo sapiens ATPase, Na+/K+ transporting, alpha 2 (+) polypeptide (ATP1A2), mRNA
NM_000701	Homo sapiens ATPase, Na+/K+ transporting, alpha 1 polypeptide (ATP1A1), mRNA
NM_000051	Homo sapiens ataxia telangiectasia mutated (includes complementation groups A, C and D) (ATM), mRNA
NM_001675	Homo sapiens activating transcription factor 4 (tax-responsive enhancer element B67) (ATF4), mRNA
NM 001673	Homo sapiens asparagine synthetase (ASNS), mRNA
NM 000048	Homo sapiens argininosuccinate lyase (ASL), mRNA
NM_001670	Homo sapiens armadillo repeat gene deletes in velocardiofacial syndrome (ARVCF), mRNA
NM 001179	Homo sapiens ADP-ribosyltransferase 3 (ART3), mRNA
NM 000047	Homo sapiens arylsulfatase E (chondrodysplasia punctata 1) (ARSE), mRNA
NM_001178	Homo sapiens aryl hydrocarbon receptor nuclear translocator-like (ARNTL), mRNA
NM 001668	Homo sapiens aryl hydrocarbon receptor nuclear translocator (ARNT), mRNA
NM 001667	Homo sapiens ADP-ribosylation factor-like 2 (ARL2), mRNA
NM_001176	Homo sapiens Rho GDP dissociation inhibitor (GDI) gamma (ARHGDIG), mRNA
NM 001665	Homo sapiens ras homolog gene family, member G (rho G) (ARHG), mRNA
NM_001661	Homo sapiens ADP-ribosylation factor 4-like (ARF4L), mRNA
NM 001659	Homo sapiens ADP-ribosylation factor 3 (ARF3), mRNA
NM_001657	Homo sapiens amphiregulin (schwannoma-derived growth factor) (AREG), mRNA
NM_001654	Homo sapiens v-raf murine sarcoma 3611 viral oncogene homolog 1 (ARAF1), mRNA
NM 001169	Homo sapiens aquaporin 8 (AQP8), mRNA
NM 001651	Homo sapiens aquaporin 5 (AQP5), mRNA
NM 001648	Homo sapiens kallikrein 3, (prostate specific antigen) (KLK3), mRNA
NM_000484	Homo sapiens amyloid beta (A4) precursor protein (protease nexin-II, Alzheimer
	disease) (APP), mRNA

NM_001647	Homo sapiens apolipoprotein D (APOD), mRNA
NM 001646	Homo sapiens apolipoprotein C-IV (APOC4), mRNA
NM 000384	Homo sapiens apolipoprotein B (including Ag(x) antigen) (APOB), mRNA
NM 001643	Homo sapiens apolipoprotein A-II (APOA2), mRNA
NM 001168	Homo sapiens baculoviral IAP repeat-containing 5 (survivin) (BIRC5), mRNA
NM 001167	Homo sapiens baculoviral IAP repeat-containing 4 (BIRC4), mRNA
NM 001164	Homo sapiens amyloid beta (A4) precursor protein-binding, family B, member 1
_	(Fe65) (APBB1), mRNA
NM_001163	Homo sapiens amyloid beta (A4) precursor protein-binding, family A, member 1
-	(X11) (APBA1), mRNA
NM 001161	Homo sapiens nudix (nucleoside diphosphate linked moiety X)-type motif 2
_	(NUDT2), mRNA
NM_001637	Homo sapiens acyloxyacyl hydrolase (neutrophil) (AOAH), mRNA
NM 001630	Homo sapiens annexin A8 (ANXA8), mRNA
NM 003568	Homo sapiens annexin A9 (ANXA9), mRNA
NM_000700	Homo sapiens annexin A1 (ANXA1), mRNA
NM_001152	Homo sapiens solute carrier family 25 (mitochondrial carrier; adenine nucleotide
	translocator), member 5 (SLC25A5), nuclear gene encoding mitochondrial
	protein, mRNA
NM_001151	Homo sapiens solute carrier family 25 (mitochondrial carrier; adenine nucleotide
	translocator), member 4 (SLC25A4), nuclear gene encoding mitochondrial
	protein, mRNA
NM_001150	Homo sapiens alanyl (membrane) aminopeptidase (aminopeptidase N,
	aminopeptidase M, microsomal aminopeptidase, CD13, p150) (ANPEP), mRNA
NM_001146	Homo sapiens angiopoietin 1 (ANGPT1), mRNA
NM_000699	Homo sapiens amylase, alpha 2A; pancreatic (AMY2A), mRNA
NM_000481	Homo sapiens aminomethyltransferase (glycine cleavage system protein T)
	(AMT), mRNA
NM_000480	Homo sapiens adenosine monophosphate deaminase (isoform E) (AMPD3),
	mRNA
NM_001144	Homo sapiens autocrine motility factor receptor (AMFR), mRNA
NM_001143	Homo sapiens amelogenin (Y chromosome) (AMELY), mRNA
NM_001633	Homo sapiens alpha-1-microglobulin/bikunin precursor (AMBP), mRNA
NM_000698	Homo sapiens arachidonate 5-lipoxygenase (ALOX5), mRNA
NM_001140	Homo sapiens arachidonate 15-lipoxygenase (ALOX15), mRNA
NM_001139	Homo sapiens arachidonate 12-lipoxygenase, 12R type (ALOX12B), mRNA
NM_000697	Homo sapiens arachidonate 12-lipoxygenase (ALOX12), mRNA
NM_001628	Homo sapiens aldo-keto reductase family 1, member B1 (aldose reductase)
	(AKR1B1), mRNA
NM_000696	Homo sapiens aldehyde dehydrogenase 9 (gamma-aminobutyraldehyde
	dehydrogenase, E3 isozyme) (ALDH9), mRNA
NM_000692	Homo sapiens aldehyde dehydrogenase 5 (ALDH5), mRNA
NM_003748	Homo sapiens aldehyde dehydrogenase 4 (glutamate gamma-semialdehyde
	dehydrogenase; pyrroline-5-carboxylate dehydrogenase) (ALDH4), mRNA
NM_000690	Homo sapiens aldehyde dehydrogenase 2, mitochondrial (ALDH2), mRNA
NM_000689	Homo sapiens aldehyde dehydrogenase 1, soluble (ALDH1), mRNA
NM_001627	Homo sapiens activated leucocyte cell adhesion molecule (ALCAM), mRNA
NM_000688	Homo sapiens aminolevulinate, delta-, synthase 1 (ALAS1), nuclear gene
37 6 600 600	encoding mitochondrial protein, mRNA
NM_003689	Homo sapiens aldo-keto reductase family 7, member A2 (aflatoxin aldehyde
37 6 000000	reductase) (AKR7A2), mRNA
NM_003886	Homo sapiens A kinase (PRKA) anchor protein 4 (AKAP4), mRNA

NM 001348 Homo sapiens A kinase (PRKA) anchor protein 1 (AKAP1), mRNA NM 001352 Homo sapiens alpha-2-Hs. glycoprotein (AHSG), mRNA NM 001131 Homo sapiens adamin (AFM), mRNA NM 001314 Homo sapiens adaptor-related protein complex 3, delta 1 subunit (AP3D1), mRNA NM 001127 Homo sapiens adaptor-related protein complex 1, beta 1 subunit (AP1B1), mRNA NM 0010127 Homo sapiens adenosine A2b receptor (ADORA2B), mRNA NM 000676 Homo sapiens adenosine A2b receptor (ADORA1B), mRNA NM 001124 Homo sapiens adenosine A1 receptor (ADORA1B), mRNA NM 001124 Homo sapiens adenosine A1 receptor (ADORA1), mRNA NM 001134 Homo sapiens adenosine A1 receptor (ADORA1B), mRNA NM 001134 Homo sapiens adenosine A1 receptor (ADORA1B), mRNA NM 001138 Homo sapiens adenosine A1 receptor (ADORA1B), mRNA NM 001118 Homo sapiens adenosine A1 receptor Like protein (TETRAN), mRNA NM 001018 Homo sapiens adenylate cyclase activating polypeptide 1 (pituitary) receptor type 1 (ADCYAP1R1), mRNA NM 001013 Homo sapiens atminoacylase 1 (ACY1), mRNA NM 001013 Homo sapiens actival (ACR), mRNA NM 001050 Homo sapiens actival (ACR), mRNA NM 001050 Homo sapiens actival (ACR), mRNA NM 001090 Homo sapiens actival (ACR), mRNA NM 001096 Homo sapiens actival (ACR) mRNA NM 001096 Homo sapiens actival (ACR) mRNA NM 001096 Homo sapiens actival (ACR) mRNA NM 001099 Homo sapiens actival (ACR) mRNA NM 001090 Homo sapiens actival (ACR) mRNA NM 001091 Homo sapiens actival (ACR) mRNA NM 001093 Homo sapiens actival (ACR) mRNA NM 001094 Homo sapiens actival (ACR) mRNA NM 001095 Homo sapiens actival (ACR) mRNA NM 001096 Homo sapiens actival (ACR) mRNA NM 001097 Homo sapiens actival (ACR) mRNA NM 001098 Homo sapiens actival (ACR) mRNA NM 00160 Homo sapiens actival (ACR) mRNA NM 00160 Homo sapie		
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	NM_001396	Homo sapiens dual-specificity tyrosine-(Y)-phosphorylation regulated kinase 1

CLAIMS

What we claim is:

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- A double-stranded short interfering nucleic acid (siNA) molecule that downregulates expression of an endogenous mammalian target gene, wherein said siNA molecule comprises one or more chemical modifications and each strand of said double-stranded siNA comprises about 21 nucleotides.
 - 2. The siNA molecule of claim 1, wherein said siNA molecule comprises no ribonucleotides.
- 3. The siNA molecule of claim 1, wherein said siNA molecule comprises ribonucleotides.
 - 4. The siNA molecule of claim 1, wherein one of the strands of said double-stranded siNA molecule comprises a nucleotide sequence that is complementary to a nucleotide sequence of the endogenous mammalian target gene or a portion thereof, and wherein the second strand of said double-stranded siNA molecule comprises a nucleotide sequence substantially similar to the nucleotide sequence of the endogenous mammalian target gene or a portion thereof.
 - 5. The siNA molecule of claim 4, wherein each strand of the siNA molecule comprises about 19 to about 23 nucleotides, and wherein each strand comprises at least about 19 nucleotides that are complementary to the nucleotides of the other strand.
 - 6. The siNA molecule of claim 1, wherein said siNA molecule comprises an antisense region comprising a nucleotide sequence that is complementary to a nucleotide sequence of the endogenous mammalian target gene or a portion thereof, and wherein said siNA further comprises a sense region, wherein said sense region comprises a nucleotide sequence substantially similar to the nucleotide sequence of said endogenous mammalian target gene or a portion thereof.
- 7. The siNA molecule of claim 6, wherein said antisense region and said sense region each comprise about 19 to about 23 nucleotides, and wherein said antisense region comprises at least about 19 nucleotides that are complementary to nucleotides of the sense region.

8. The siNA molecule of claim 1, wherein said siNA molecule comprises a sense region and an antisense region and wherein said antisense region comprises a nucleotide sequence that is complementary to a nucleotide sequence of RNA encoded by the endogenous mammalian target gene or a portion thereof and said sense region comprises a nucleotide sequence that is complementary to said antisense region.

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- 9. The siNA molecule of claim 6, wherein said siNA molecule is assembled from two separate oligonucleotide fragments, wherein one fragment comprises the sense region and the second fragment comprises the antisense region of said siNA molecule.
- 10. The siNA molecule of claim claim 6, wherein said sense region is connected to the antisense region via a linker molecule.
- 11. The siNA molecule of claim 10, wherein said linker molecule is a polynucleotide linker.
- 15 12. The siNA molecule of claim 10, wherein said linker molecule is a non-nucleotide linker.
 - 13. The siNA molecule of claim 6, wherein pyrimidine nucleotides in the sense region are 2'-O-methyl pyrimidine nucleotides.
- 14. The siNA molecule of claim 6, wherein purine nucleotides in the sense region are 2'-deoxy purine nucleotides.
 - 15. The siNA molecule of claim 6, wherein the pyrimidine nucleotides present in the sense region are 2'-deoxy-2'-fluoro pyrimidine nucleotides.
 - 16. The siNA molecule of claim 9, wherein the fragment comprising said sense region includes a terminal cap moiety at the 5'-end, the 3'-end, or both of the 5' and 3' ends of the fragment comprising said sense region.
 - 17. The siNA molecule of claim 16, wherein said terminal cap moiety is an inverted deoxy abasic moiety.
 - 18. The siNA molecule of claim 6, wherein the pyrimidine nucleotides of said antisense region are 2'-deoxy-2'-fluoro pyrimidine nucleotides.

19. The siNA molecule of claim 6, wherein the the purine nucleotides of said antisense region are 2'-O-methyl purine nucleotides.

- 20. The siNA molecule of claim 6, wherein the purine nucleotides present in said antisense region comprise 2'-deoxy- purine nucleotides.
- 5 21. The siNA molecule of claim 18, wherein said antisense region comprises a phosphorothicate internucleotide linkage at the 3' end of said antisense region.
 - 22. The siNA molecule of claim 6, wherein said antisense region comprises a glyceryl modification at the 3' end of said antisense region.
- The siNA molecule of claim 9, wherein each of the two fragments of said siNA molecule comprise 21 nucleotides.
 - 24. The siNA molecule of claim 23, wherein about 19 nucleotides of each fragment of the siNA molecule are base-paired to the complementary nucleotides of the other fragment of the siNA molecule and wherein at least two 3' terminal nucleotides of each fragment of the siNA molecule are not base-paired to the nucleotides of the other fragment of the siNA molecule.

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- 25. The siNA molecule of claim 24, wherein each of the two 3' terminal nucleotides of each fragment of the siNA molecule are 2'-deoxy-pyrimidines.
- 26. The siNA molecule of claim 25, wherein said 2'-deoxy-pyrimidine is 2'-deoxy-thymidine.
- 20 27. The siNA molecule of claim 23, wherein all 21 nucleotides of each fragment of the siNA molecule are base-paired to the complementary nucleotides of the other fragment of the siNA molecule.
 - 28. The siNA molecule of claim 23, wherein about 19 nucleotides of the antisense region are base-paired to the nucleotide sequence of the RNA encoded by the endogenous mammalian target gene or a portion thereof.
 - 29. The siNA molecule of claim 23, wherein 21 nucleotides of the antisense region are base-paired to the nucleotide sequence of the RNA encoded by the endogenous mammalian target gene or a portion thereof.
- 30. The siNA molecule of claim 9, wherein the 5'-end of the fragment comprising said antisense region optionally includes a phosphate group.

31. The siNA molecule of claim 1, wherein said mammalian gene is a human gene.

- 32. A double-stranded short interfering nucleic acid (siNA) molecule that inhibits the expression of an endogenous mammalian target RNA sequence, wherein each strand of said double-stranded siNA molecule comprises about 21 nucleotides and wherein said siNA molecule comprises no ribonucleotides.
- 33. The siNA molecule of claim 32, wherein said target RNA sequence is encoded by a human gene.
- A double-stranded short interfering nucleic acid (siNA) molecule that inhibits the expression of an endogenous mammalian target gene, wherein each strand of said double-stranded siNA molecule comprises about 21 nucleotides and wherein said siNA molecule does not require the presence of a ribonucleotide within the siNA molecule for the inhibition of expression of an endogenous mammalian target gene.
- 35. The siNA molecule of claim 34, wherein said mammalian target gene is a human gene.
 - 36. The siNA molecule of claim 31 or claim 35, wherein said human gene is vascular endothelial growth factor (VEGF).
 - 37. The siNA molecule of claim 31 or claim 35, wherein said human gene is a receptor for VEGF.
- 20 38. The siNA of claim 37, wherein said receptor is VEGFR1.

- 39. The siNA of claim 37, wherein said receptor is VEGFR2.
- 40. The siNA of claim 37, wherein said receptor is VEGFR3
- 41. The siNA molecule of claim 31 or claim 35, wherein said human gene is BCL2.
- 42. The siNA molecule of claim 31 or claim 35, wherein said human gene is HER2/neu.
 - 43. The siNA molecule of claim 31 or claim 35, wherein said human gene is c-Myc.
 - 44. The siNA molecule of claim 31 or claim 35, wherein said human gene is PCNA.
 - 45. The siNA molecule of claim 31 or claim 35, wherein said human gene is REL-A.

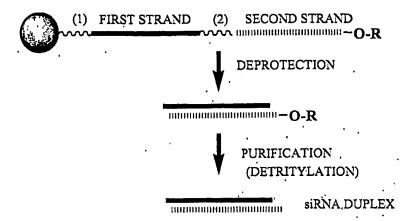
46. The siNA molecule of claim 31 or claim 35, wherein said human gene is PTP1B.

- 47. The siNA molecule of claim 31 or claim 35, wherein said human gene is BACE.
- 48. The siNA molecule of claim 31 or claim 35, wherein said human gene is CHK1.
- 49. The siNA molecule of claim 31 or claim 35, wherein said human gene is PKCalpha.
 - 50. The siNA molecule of claim 31 or claim 35, wherein said human gene is EGFR (HER1).
 - 51. A pharmaceutical composition comprising the siNA molecule of claim 1 in an acceptable carrier or diluent.
- 10 52. Medicament comprising the siNA molecule of claim 1.

- 53. Active ingredient comprising the siNA molecule of claim 1.
- 54. Use of a double-stranded short interfering nucleic acid (siNA) molecule to down-regulate expression of an endogenous mammalian target gene, wherein said siNA molecule comprises one or more chemical modifications and each strand of said double-stranded siNA comprises about 21 nucleotides.

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Figure 1



= SOLID SUPPORT

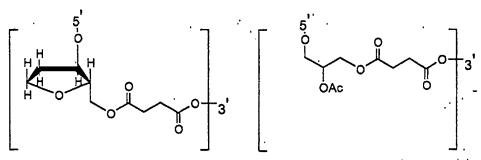
R = TERMINAL PROTECTING GROUP FOR EXAMPLE: DIMETHOXYTRITYL (DMT)

= CLEAVABLE LINKER

(FOR EXAMPLE: NUCLEOTIDE SUCCINATE OR

(2) INVERTED DEOXYABASIC SUCCINATE)

(FOR EXAMPLE: NUCLEOTIDE SUCCINATE OR INVERTED DEOXYABASIC SUCCINATE)

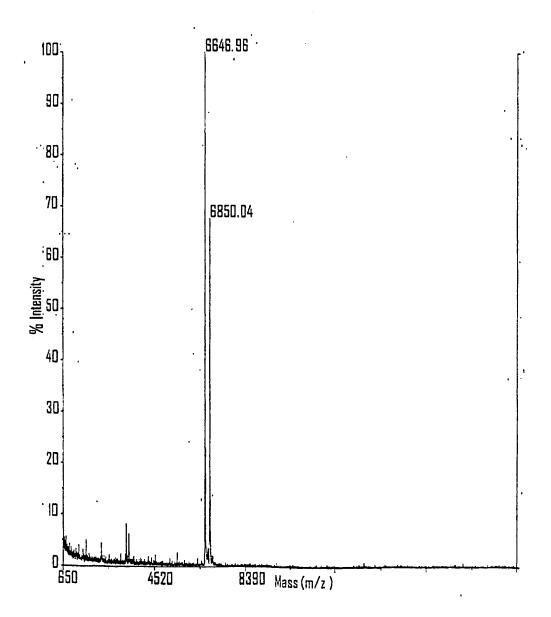


INVERTED DEOXYABASIC SUCCINATE LINKAGE

GLYCERYL SUCCINATE LINKAGE

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Figure 2



WO 03/074654

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Figure 3

5'-CGUACGCGGAAUACUUCGATT (SEQ ID NO: 925) 3'-TTGCAUGCGCCUUAUGAAGCU (SEQ ID NO: 926)

T % = 138 min5'-B cAAccACAAAuAcAACAATT B (SEQ ID NO: 925) 3'-TXGuuGGuGuuuuAuGuuGuu (SEQ ID NO: 927)

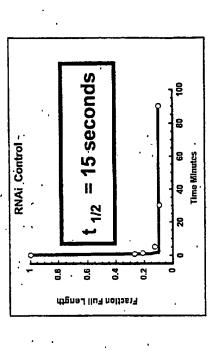
5'-B cAAccACAAAUACAACAATT B (SEQ ID NO: 925) T % = 3.7 days

3'-TDGuuGGuGuuuuAuGuuGuu (SEQ ID NO: 928)

5'-B cAAccACAAAuAcAATT B (SEQ ID NO: 925) T %=72 minutes 3'-XTGuuGGuGuuuuAuGuuGuu (SEQ ID NO: 929)

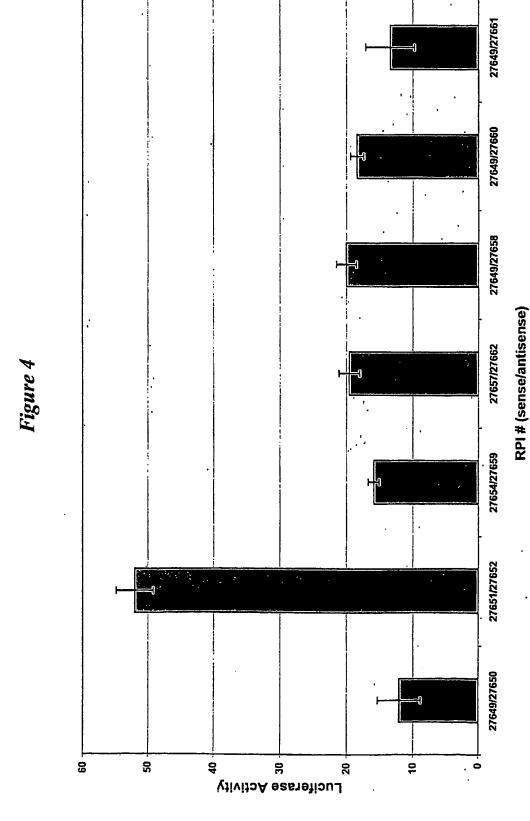
5'-B cAAccACAAAAuAcAACTT B (SEQ ID NO: 925) T $\frac{1}{2}$ = 40 days 3'-LTGuuGGuGuuuuAuGuuGuu (SEQ ID NO: 930)

5'-B cAAccACAAAUACAACAATT B (SEQ ID NO: 925) $T \frac{1}{2} = 32 \text{ days}$ 3'-tTGuuGGuGuuuuAuGuuGuu (SEQ ID NO: 931)

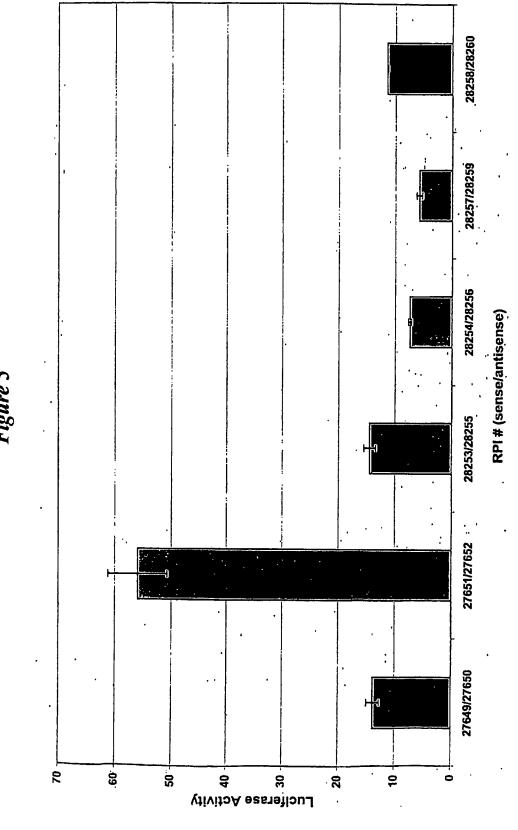


G, A, U, C = Guanosine, Adenosine, Uridine, Cytidine Lower Case = 2'-deoxy-2'-fluoro B = inverted deoxyabasic X = 3'-deoxy Thymidine D = inverted Thymidine S = phosphorothioate G = terminal glycine = L-thymidine T = Thymidine

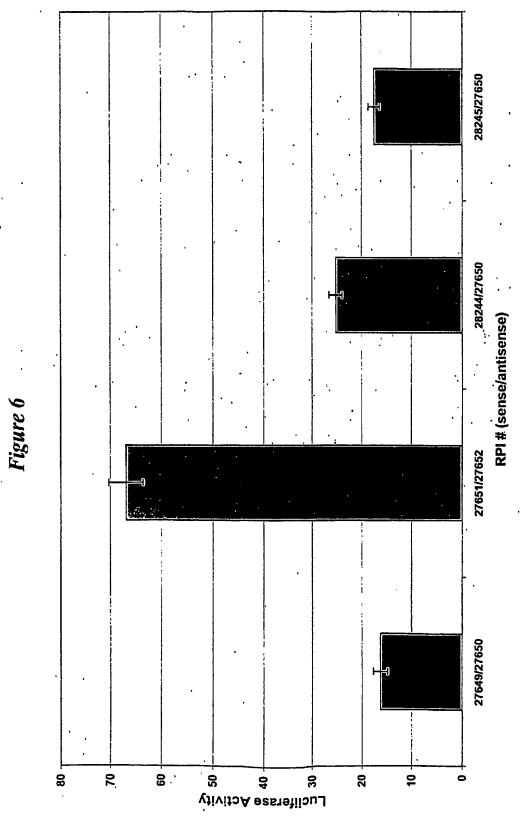
L = Glyceryl moiety



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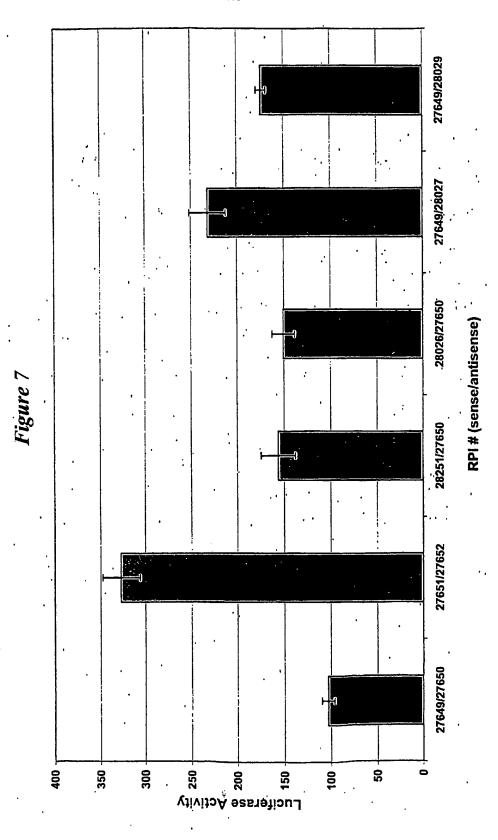


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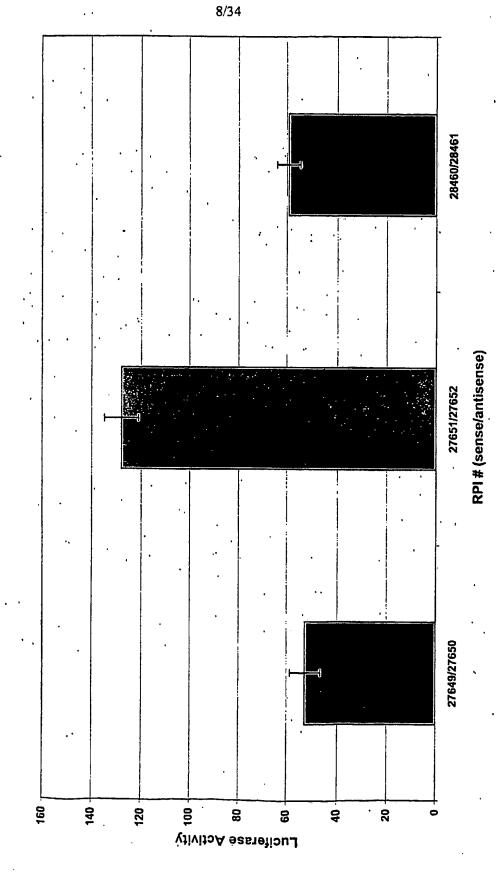


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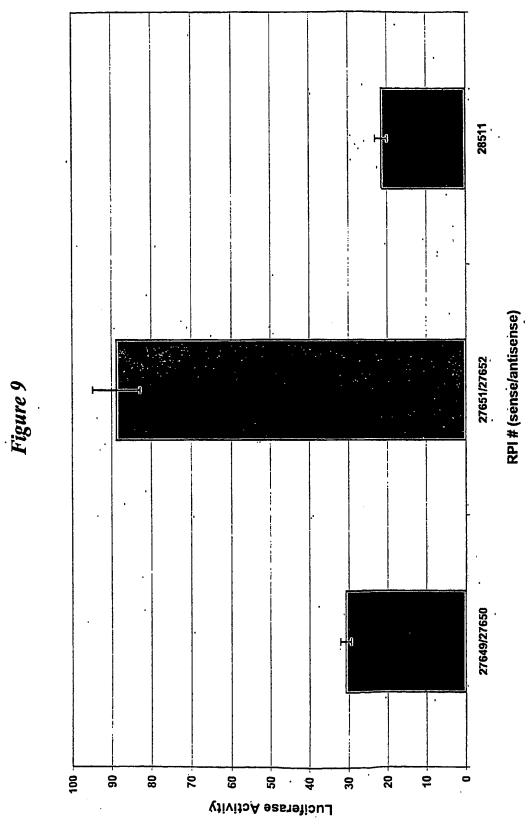




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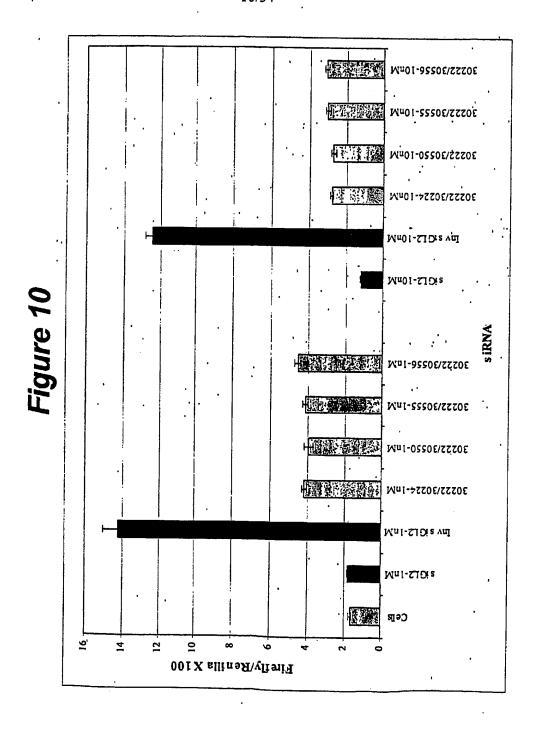


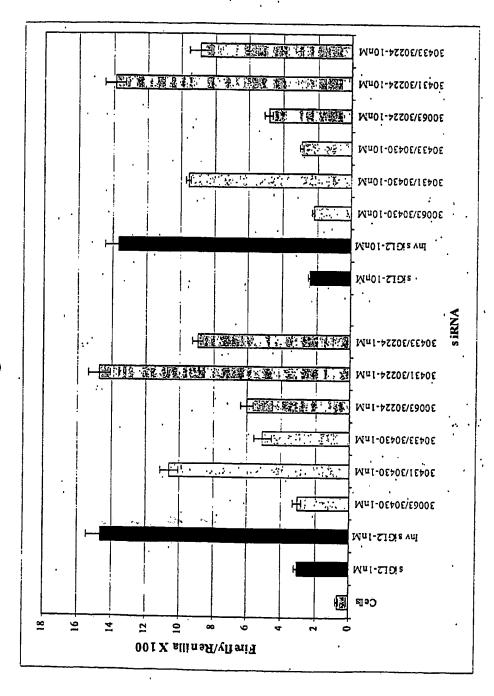
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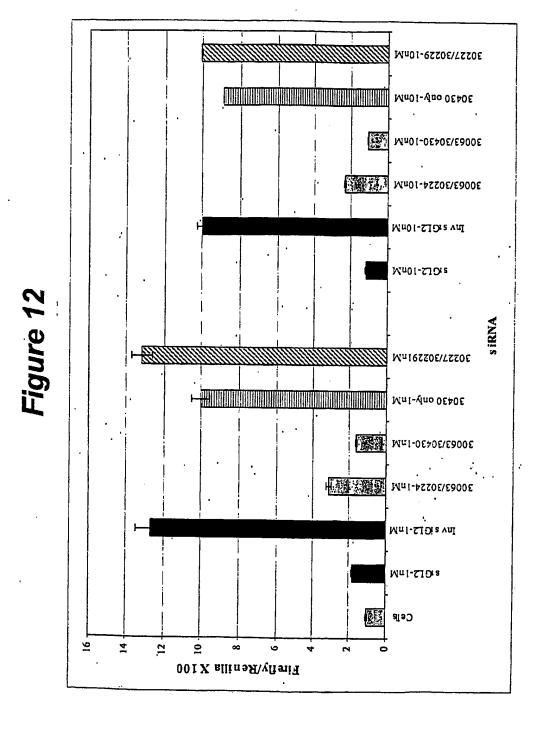




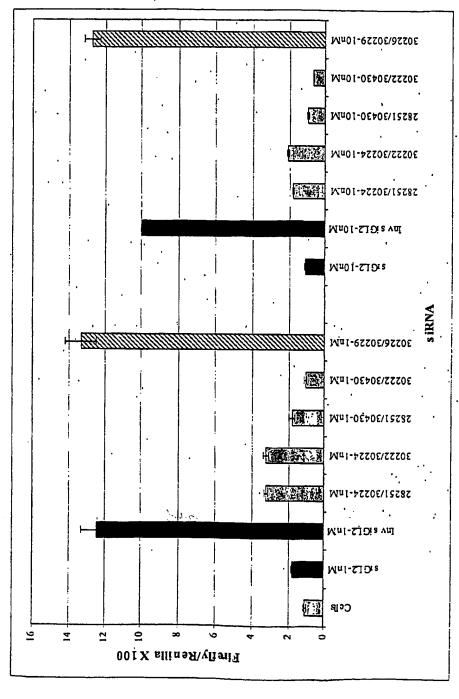


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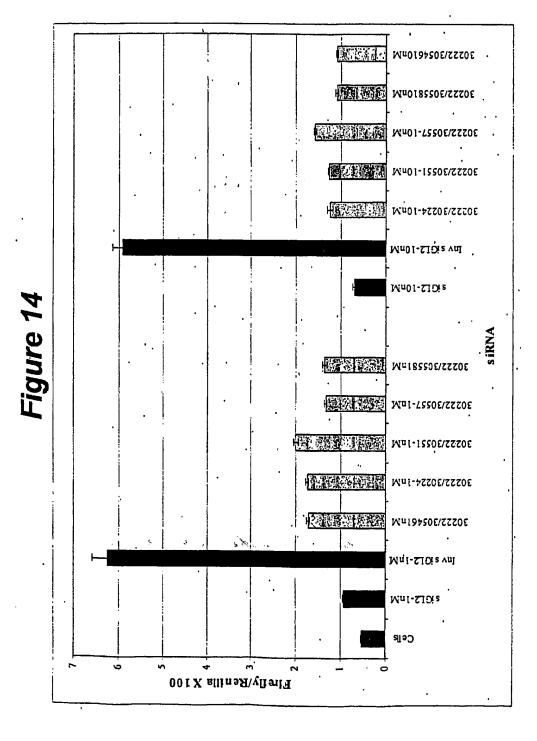




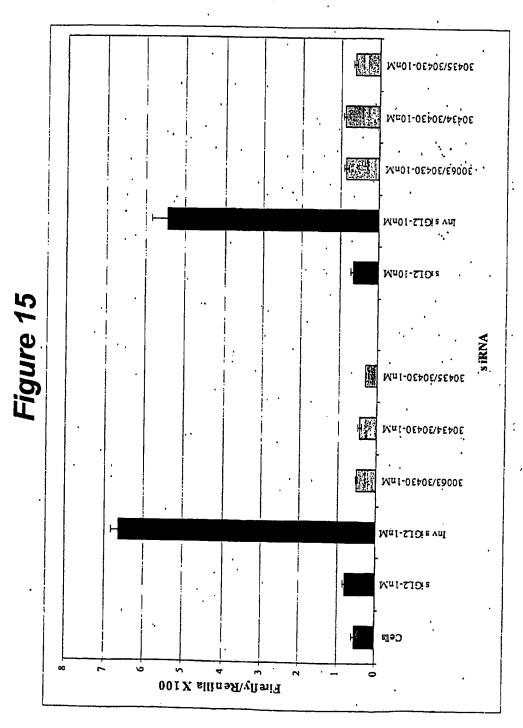


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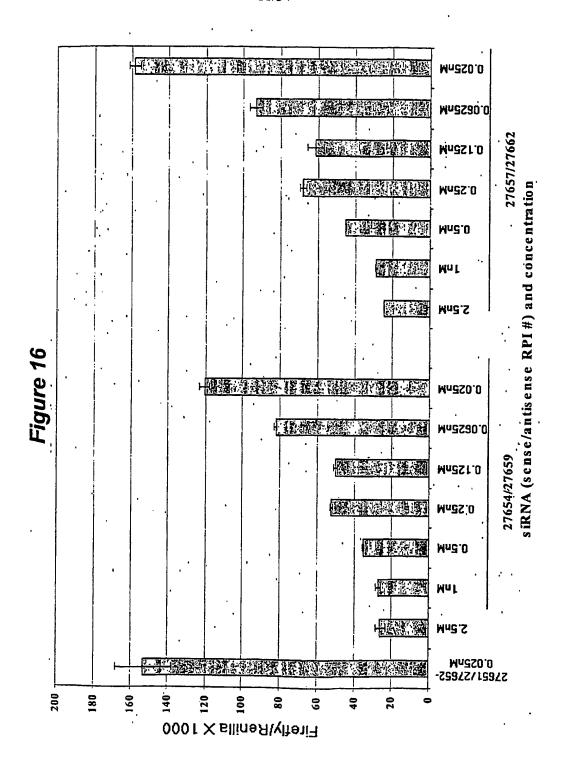




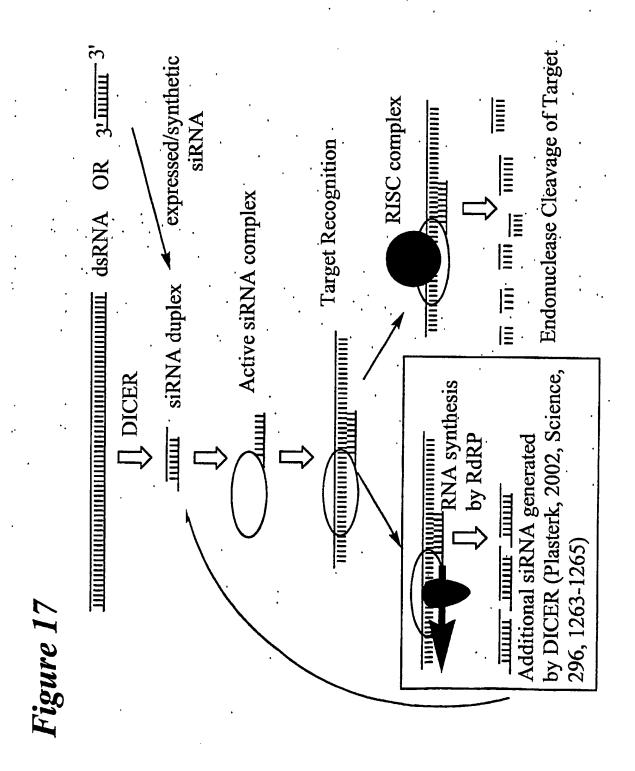




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Figure 18 SENSE STRAND (SEQ ID NO 903) ALL PYRIMIDINES = 2'-O-ME OR 2'-FLUORO EXCEPT POSITIONS (N N) -3' 3'- $L-(N_sN)NNNNNNNNNNNNNNNNNSN_sN_sN_sN_sN_s$ -5' ANTISENSE STRAND (SEQ ID NO 904) ALL PYRIMIDINES = 2'-FLUORO EXCEPT POSITIONS (N N) SENSE STRAND (SEQ ID NO 905) ALL PYRIMIDINES = 2'-O-ME OR 2'-FLUORO EXCEPT POSITIONS (N N) ·-3' B 3'--5' ANTISENSE STRAND (SEQ ID NO 906) ALL PYRIMIDINES = 2'-FLUORO EXCEPT POSITIONS (N:N) SENSE STRAND (SEQ ID NO 907) ALL PYRIMIDINES = 2'-O-ME OR 2'-FLUORO EXCEPT POSITIONS (N N) -3' 3'--5' ANTISENSE STRAND, (SEQ ID NO 908) ALL PYRIMIDINES = 2'-FLUORO EXCEPT POSITIONS (N N) SENSE STRAND (SEQ ID NO 909) ALL PYRIMIDINES = 2'-FLUORO EXCEPT POSITIONS (N N) AND ALL PURINES = 2'-DEOXY 5'-B-NNNNNNNNNNNNNNNNNNNNNNNNNNNN -3' 3'--5' ANTISENSE STRAND (SEQ ID NO 910) ALL PYRIMIDINES = 2'-FLUORO AND ALL PURINES = 2'-O-ME EXCEPT POSITIONS (N N) SENSE STRAND (SEQ ID NO 911) ALL PYRIMIDINES = 2'-FLUORO EXCEPT POSITIONS (N N) -3' \mathbf{E} L-(NN) NNNNNNNNNNNNNNNNNNN ANTISENSE STRAND (SEO ID NO 912) ALL PYRIMIDINES = 2'-FLUORO AND ALL PURINES = 2'-O-ME EXCEPT POSITIONS (N N) : SENSE STRAND (SEQ ID NO 909) ALL PYRIMIDINES = 2'-FLUORO EXCEPT POSITIONS (N N) AND ALL PURINES = 2'-DEOXY 5'--3' F 3'--5' ANTISENSE STRAND (SEQ ID NO 913) ALL PYRIMIDINES = 2'-FLUORO EXCEPT POSITIONS (N N) AND ALL PURINES = 2'-DEOXY

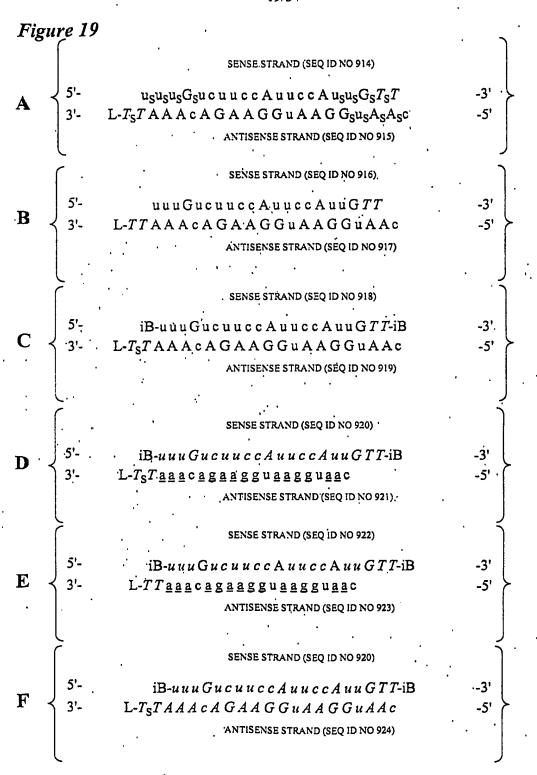
POSITIONS (NN) CAN COMPRISE ANY NUCLEOTIDE, SUCH AS DEOXYNUCLEOTIDES (eg. THYMIDINE) OR UNIVERSAL BASES

B = ABASIC, INVERTED ABASIC, INVERTED NUCLEOTIDE OR OTHER TERMINAL CAP THAT IS OPTIONALLY PRESENT

L = GLYCERYL MOIETY THAT IS OPTIONALLY PRESENT

S = PHOSPHOROTHIOATE OR PHOSPHORODITHIOATE

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lower case = 2'-O-Methyl or 2'-deoxy-2'-fluoro; italic lower case = 2'-deoxy-2'-fluoro ITALIC UPPER CASE = DEOXY B = INVERTED DEOXYABASIC
L = GLYCERYL MOIETY OPTIONALLY PRESENT
S = PHOSPHOROTHIOATE OR
PHOSPHORODITHIOATE



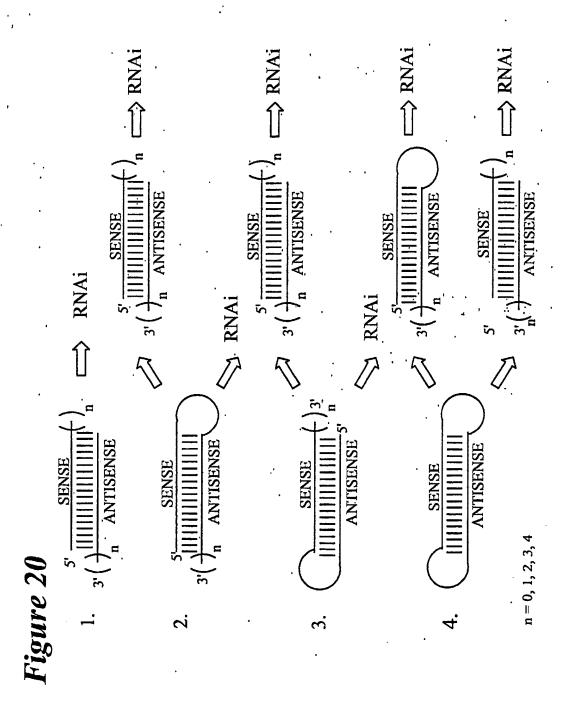
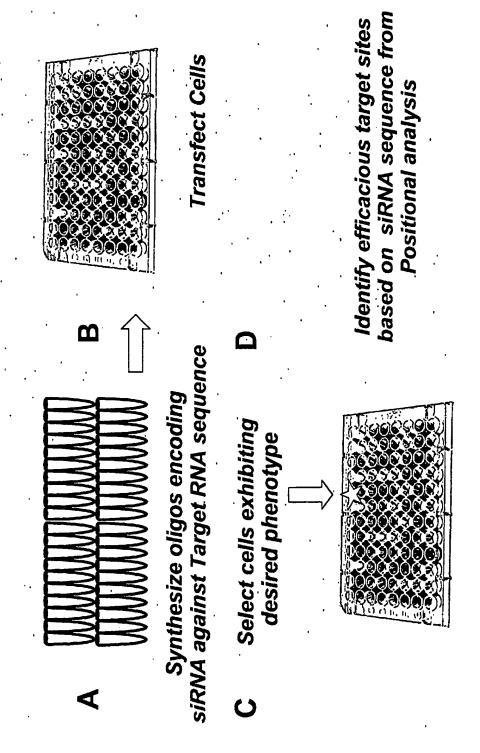


Figure 21: Target site Selection using sik



R = O, S, N, alkyl, substituted alkyl, O-alkyl, S-alkyl, alkaryl, or aralkyl B = Independently any nucleotide base, either naturally occurring or chemically modified, or optionally H (abasic).

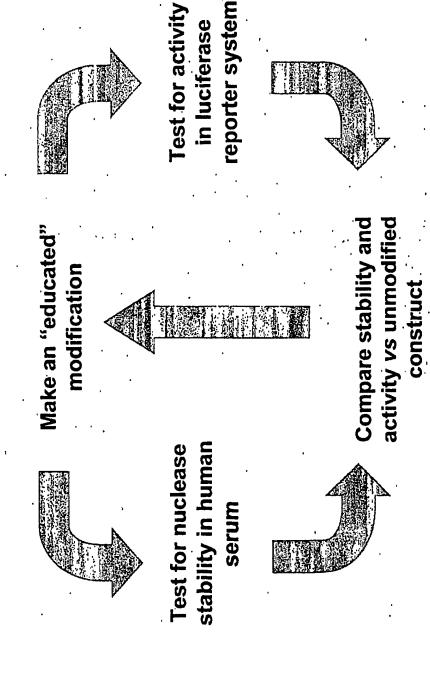
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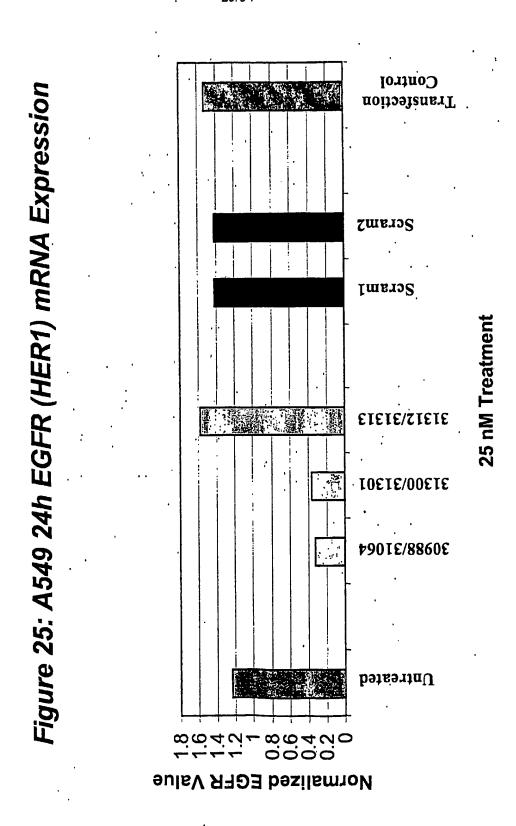
VEGF Control Figure 23: Inhibition of VEGF-Induced Angiogenesis API 29983/29984 (nactive·lug *p< 0.05 with respect to VEGF by Dunnett's
**p<0.05 with respect to Inverted control by Tukcy-Kramer by siRNAs RPI 29983/29984 RPI 29695/29699 Active 1úg. KPI 29695/29699. Active 10ug RPI 29695/29699 100 80 20 9 5 0

b oubrii TOEGF induc d % sizenegoignA

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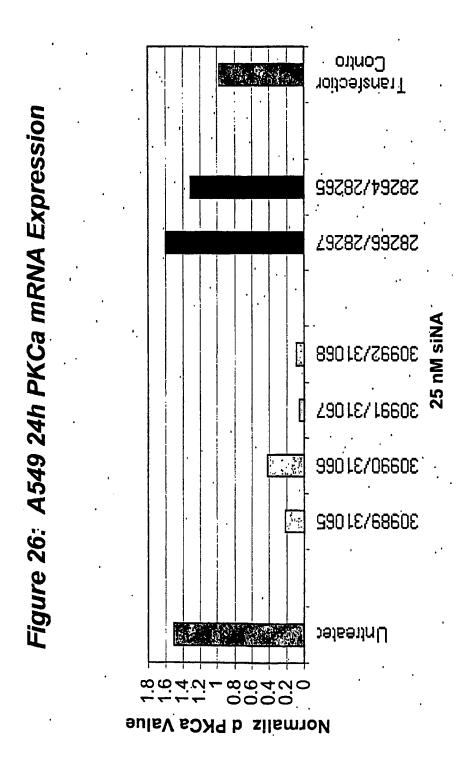






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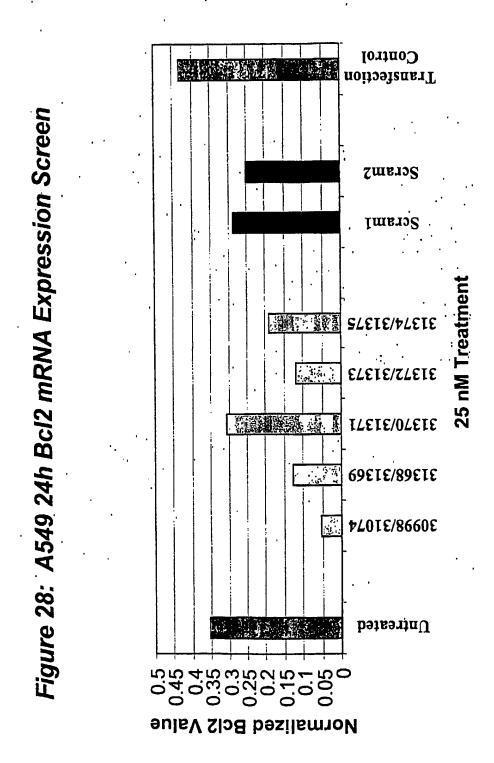
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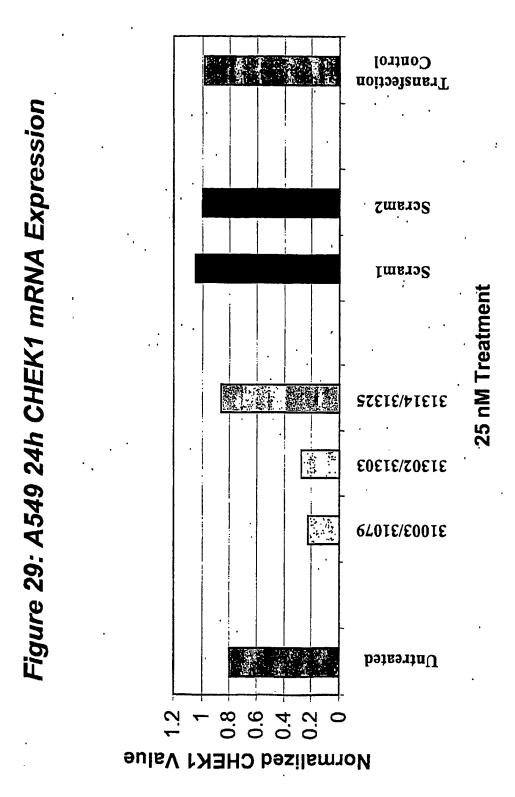
Control Transfection · Figure 27: siNA mediated inhibition of MYC RNA Seram Seraml. ZL01E/9660E //... 1/016/56608 30994/31072 6901E/E660E Untreated 0.25 0.2 0.15 0.1 Normalized MYC Value

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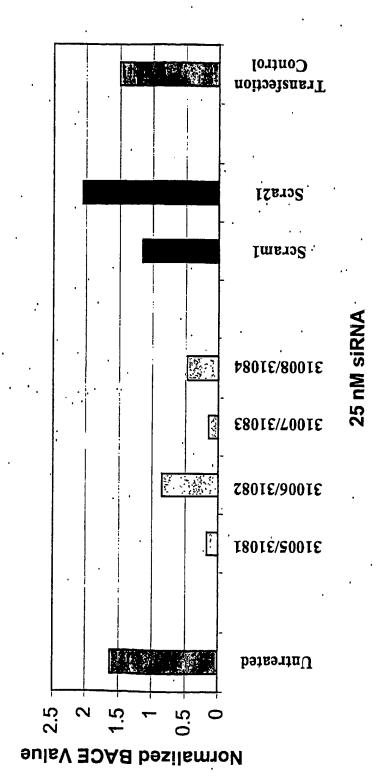
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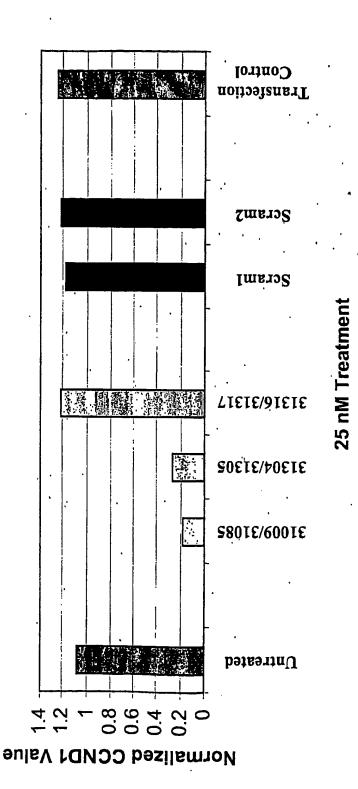
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Figure 30: A549 24h BACE mRNA Expression



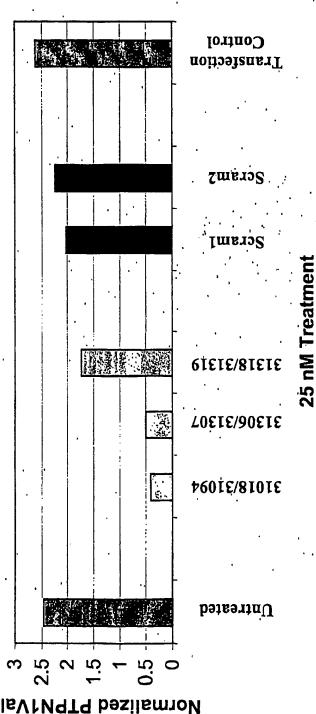
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Figure 31: A549 24h CCND1 mRNA Expression

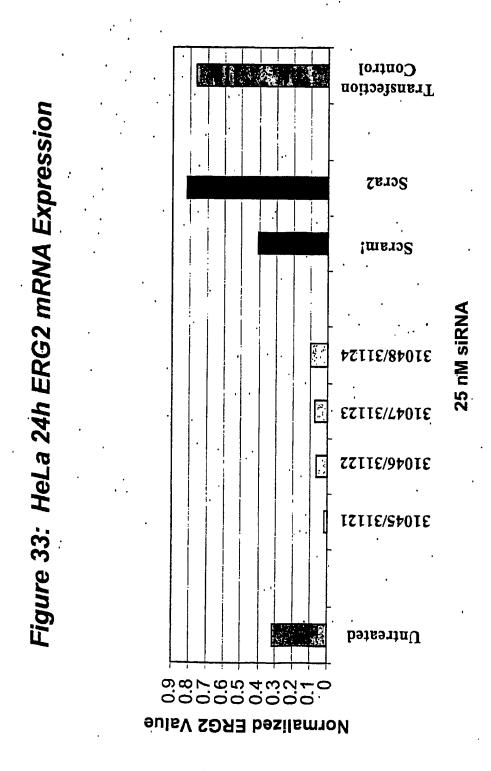


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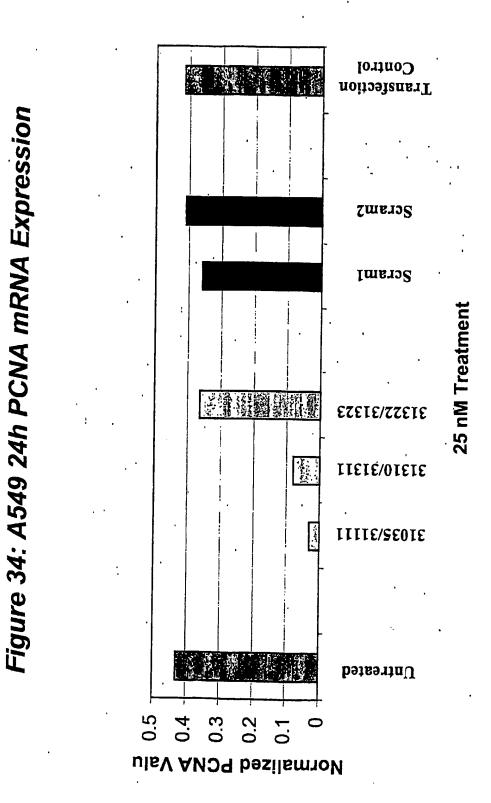


Normalized PTPN1Value



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